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60 57.1 18 4 US-09-081-975-22 Sequence 22, 59 56.2 11 1 US-08-277-660A-9 Sequence 9, 59 56.2 11 1 US-08-424-957-17 Sequence 17, 59 56.2 11 4 US-09-035-686-17 Sequence 17,	23	65	61.9	15	ß	PCT-US95-02856-6	9
59 56.2 11 1 US-08-277-660A-9 Sequence 9, 59 56.2 11 1 US-08-424-957-17 Sequence 17, 59 56.2 11 4 US-09-035-686-17 Sequence 17,	24	9	57.1	18	4	US-09-081-975-22	22
59 56.2 11 1 US-08-424-957-17 Sequence 17, 59 56.2 11 4 US-09-035-686-17 Sequence 17,	25	29	56.2	11	-	US-08-277-660A-9	
59 56.2 11 4 US-09-035-686-17 Sequence 17,	56	59	9	11	٦	US-08-424-957-17	17
	27	59	56.2	11	4	US-09-035-686-17	17,

Sequence 5, Appli Sequence 45, Appl Sequence 46, Appl Sequence 46, Appl Sequence 7, Appl Sequence 7, Appli Sequence 10, Appli Sequence 23, Appl Sequence 28, Appl	THERETO, AND USES OF THE	0; s 0; Gaps 0;
4 US-09-236-415-5 1 US-08-424-957-46 4 US-09-035-686-45 4 US-09-035-686-46 4 US-09-400-653A-7 4 US-09-400-653A-7 1 US-08-277-660A-15 1 US-08-277-660A-15 1 US-08-277-660A-15 1 US-08-424-957-23 1 US-08-424-957-28 1 US-09-035-686-23 4 US-09-035-686-24 4 US-09-035-686-40 1 US-08-424-957-11 1 US-08-424-957-11	ALIGNMENTS S TREATMENT USING IBODIES AND PROTEINS 1,975 1,975	94.3%; Score 99; DB 4; Length 20 100.0%; Pred. No. 1.6e-09; iive 0; Mismatches 0; Indels
28 59 56.2 113 23 57 54.3 111 32 57 54.3 111 32 33 57 54.3 111 33 57 54.3 111 35 56 53.3 111 38 56 53.3 111 55 55 52.4 100	RESULT 1 US-09-081-975-8 Sequence 8, Application US/09081975 Patent No. 6451979 GENERAL INFORMATION: APPLICANT: Kaelin, William APPLICANT: Aselin, WILLEOF INVENTION: METHODS OF TREATMENT ITILE OF INVENTION: METHODS OF TREATMENT ITILE OF INVENTION: ANTIBODIES NUMBER OF SEQUENCES: 27 CORRESPONDENCE ADDRESS: ADDRESSE: Nixon Peabody LLP STREET: 101 Federal Street CITY: Boston STATE: MA COUNTRY: USA ZIP: OCLINO STATE: MA COUNTRY: USA ZIP: OCHENENT APPLICATION DATA: MEDIUM TYPE: Diskette COMPUTER: EsastsEQ for Windows SOFTWARE: FastsEQ for Windows SOFTWARE: L2-MAY-1998 CLASSIFICATION NUMBER: US/09/081,975 FILING DATE: 12-MAY-1997 FILING DATE: 13-MAY-1997 FILING DATE: 12-MAY-1997 FILING DATE: 13-MAY-1997 F	Query Match Best Local Similarity 10 Matches 18; Conservativ

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RESULT 4
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Patent No. 6451979

GENERAL INFORMATION:

APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCES: 27
CORRESPONDENCES: 27
CORRESPONDENCES: 27
CORRESPONDENCES: 27
                                                                                                                                                                                                              APPLICANT: Xeelin, William
APPLICANT: Xeelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHOD SOF TREATMENT USING
TITLE OF INVENTION: MATIBODIES
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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94.3%; Score 99; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.6e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION DATA:
APPLICATION MUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                       Sequence 10, Application US/09081975
Patent No. 6451979
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 4740
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nixon Peabody LLP
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                     2 PPLSQETFSDLWKLLPEN 19
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1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                              Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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                                                                                                                                 US-09-081-975-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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US-09-081-975-16

Sequence 16, Application US/09081975

Patent No. 6451979

GENERAL INFORMATION:
APPLICANT: Acelin, William
APPLICANT: JOSE, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
TUTLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.3%; Score 99; DB 4; Length 20; 100.0%; Pred. No. 1.6e-09; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
                                                                                                                                              COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 45
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Rohald I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 30,628
REPERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  E: Nixon Peabody LLP
101 Federal Street
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101 Federal Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 100.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
                                                                                                                                  Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: linear
US-09-081-975-11
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                         Boston
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ZIP: 02110
                                                                                    02110
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                                                                 COUNTRY:
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  STREET:
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APPLICANT: Kaelin, William
APPLICANT: JOSE, Christine
APPLICANT: JOSE, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
WUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 13, Application US/09081975
Patent No. 6451979
GENERAL INFORMATION:
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: FREESEQ for Windows Version 2.0b CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFTATION: 435
PRIOR APPLICATION NATA:
APPLICATION NUMBER: 60/046,207
FILING APPLICATION SATA:
APPLICATION NUMBER: 60/046,207
ATTORNEY AGENT INFORMATION:
NAME: Bisenstein, Ronald I
                                                                                                                                                                                             Sequence 15, Application US/09081975; Patent No. 6451979; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM Compatible OPERATING SYSTEM: Windows
                                      1 PPLSOETFSDLWKLLPEN 18
                                                                   2 PPLSQETFSDLWNLLPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 PPLSQETFSDLWNLLPEN 19
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Best Local Similarity 94.4'
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 617-345-1300
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: linear
US-09-081-975-15
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STREET: 101
TW: Boston
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US-09-081-975-13
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Patent No. 6451979
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCES 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: ALXON Peabody LLP
STREET: 101 Federal Street
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94.3%; Score 99; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.6e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
89.5%; Score 94; DB 4; Length 20;
Best Local Similarity 94.4%; Pred. No. 1e-08;
Matches 17; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IRM Compatible
COMPUTER: IRM Compatible
OPERATING SYSTEM: Windows
SOFTWAKE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY-AGENT INFORMATION:
which contain Data:
ATTORNEY-AGENT INFORMATION:
which contains Data:
ATTORNEY-AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEPRAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 47400
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           30,628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 14: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                           1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                       INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 amino acids
                                                                                                                                                                                                                               ; TYPE: amino acid
; STRANDEDNESS: single
; TOPOLOGY: linear
US-09-081-975-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid STRANDEDNESS: si
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: US.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-081-975-14
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APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHOD SOF TREATMENT USING
TITLE OF INVENTION: MATIBODIES
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LiP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ô
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85.7%; Score 90; DB 4; Length 19;
Best Local Similarity 88.9%; Pred. No. 4.2e-08;
Matches 16; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: Fast SEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
                                                                                                                                                     MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FeatCSD for Windows
SOFTWARE: FeatCSD for Windows
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION 13-MAY-1998
REDICATION DATE: 60/046,207
FILING DATE: 12-MAY-1997
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 19, Application US/09081975 Patent No. 6451979
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       47400
                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFRERUCE/DOCKET NUMBER: 47401
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEPAX: 617-345-1300
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    13:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            single
                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ropology: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
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STREET: 10-
TTY: Boston
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                                                                                       USA
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                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Lane, David P.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS: ADDRESSEE: Plehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
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                                                                                                                                                                                                                                                                                                                                            81.9%; Score 86; DB 4; Length 20; 94.1%; Pred. No. 1.9e-07; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUBER: US/08/277,660A FILING DATE: 20-JUL-1994 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 73.3%; Score 77; DB 1; Le Best Local Similarity 100.0%; Pred. No. 3.8e-06; Matches 14; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REPERNCE/DOCKET NUMBER: A-60244/WHD
TELECOMMINICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM SENDING WEDLUM TYPE: FLOPPY disk COMPUTER: IEM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                  NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFRAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-277-660A-1; Sequence 1, Application US/08277660A; Patent No. 5702908
                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS: LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Four Embarcadero
CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LECORNOL:
TELEPHONE: (415) , ...
"TIREAX: (415) 398-3249
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                            16; Conservative
                                                                                                                                                                                                                                                              single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
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US-09-081-975-19
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RESULT 12
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                                                                                                                                 Sequence 4, Application US/08277660A
| Patent No. 5702908
| GENERAL INFORMATION:
| APPLICANT: Line, David P. |
| TITLE OF INVENTION: Interruption of Binding of MDM2 and P53 |
| TITLE OF INVENTION: Interruption of Binding of MDM2 and P53 |
| TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 27 CORRESPONDENCE ADDRESS: |
| ADDRESSED: Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400 |
| CITY: San Francisco |
| STREET: California |
| COUNTRY: United States |
| COUNTRY: United States |
| CITY: San Prancisco |
| STATE: California |
| COUNTRY: United States |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-424-957-1

Sequence 1, Application US/08424957

Patent No. 2770377

GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400

CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        73.3%; Score 77; DB 1; Length 15; 100.0%; Pred. No. 3.8e-06; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
FILING DATE: 20-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: 460244/WHD
TELECOMMUNICATION INFORMATION:
TELEPHAN: (415) 78-1989
TELERY: (415) 78-1089
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 PPLSQETFSDLWKL 15
1 QETFSDLWKLLPEN 14
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
                                                                                 RESULT 10
US-08-277-660A-4
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COMPERATING SYSTEM: INPR Comparations
SOUTHWER: INPR COMPETED:
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APPLICANT: Picksley, Steven M.
APPLICANT: David P.
TILLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
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                                                                                                                                                    Length 15;
                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400 CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                  Query Match 73.3%; Score 77; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.8e-06;
Matches 14; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A-61228/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-UUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-612
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 1, Application US/09035686
; Patent No. 6153391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Dreger, Walter H. REGISTRATION NUMBER: 24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RY: United States 94111-4187
                  LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 QETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                            1 PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                      2 PPLSQETFSDLWKL 15
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                                                                                       unknown
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE
                                                                                     ; TOPOLOGY:
US-08-424-957-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-035-686-1
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RESULT 14

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Gaps
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Patent No. 5702908

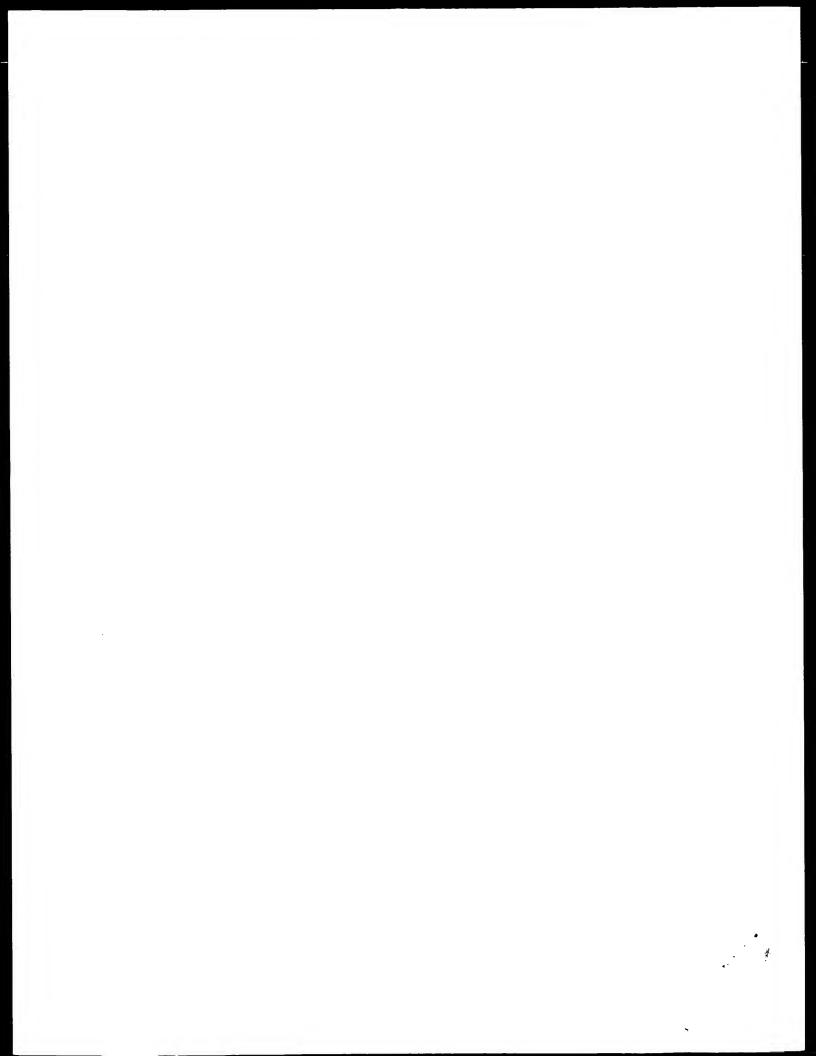
APELICANT: Pickeley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                      APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
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                                                                                                                                                                                           ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400 CITY: San Francisco STATE: California CANURY: United States ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                              PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
          ; Sequence 20, Application US/09035686; Patent No. 6153391; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEX: (415, 25.
TELEX: 910 277299
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TVPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Dreger, Walter H. REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-6
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (415) , ...
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California
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
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US-08-277-660A-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-035-686-20
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US-09-035-686-20
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STATE:
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WEDIUM TYPE: Floppy disk
COMPUTER: INN PC compatible
OPERATING SYSTEM.
COMPUTER: INN PC compatible
OPERATING SYSTEM.
CURRENT APPLICATION DATE:
PLOS/MS-DOS

FILING DATE:
CLASSIFICATION NUMBER: US/08/277,660A

FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Dreager, Malter H.
REGISTRATION INFORMATION:
FELEXON NUMBER: 24,19

REFERENCE/DOCKET NUMBER: 24
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 11:14:12; Search time 26.9167 Seconds Run on:

(without alignments)
59.144 Million cell updates/sec

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19 Perfect score:

Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

425801 segs, 83787487 residues Searched: Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Pending Patents AA New:*

1. /cgn2 6/ptodata/1/paa/PCT NEW COMB.pep:*

2. /cgn2 6/ptodata/1/paa/USO6 NEW COMB.pep:*

3. /cgn2 6/ptodata/1/paa/USO7 NEW COMB.pep:*

4. /cgn2 6/ptodata/1/paa/USO8 NEW COMB.pep:*

5. /cgn2 6/ptodata/1/paa/USO8 NEW COMB.pep:*

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6. /cgn2 6/ptodata/1/paa/USO8 NEW COMB.pep:*

7. /cgn2 6/ptodata/1/paa/USO8 NEW COMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Sequence 54, Appl Sequence 6, Appli	Sequence 149, App	Sequence 3, Appli	297,	Sequence 219, App	9		Sequence 25751, A	Sequence 29042, A	Sequence 25751, A	Sequence 29042, A	Sequence 67, Appl	18,	67,	Sequence 463, App
US-10-319-786-54 US-10-190-866A-6	US-10-283-599-149 US-09-632-036E-11	US-60-438-805-3	US-60-427-045-297	US-10-283-599-219 US-10-232-410-9	US-10-299-867-60	US-10-203-138A-15194	US-09-641-528-25751	US-09-641-528-29042	US-09-641-528A-25751	US-09-641-528A-29042	US-10-029-206A-67	US-10-301-644-18	US-10-028-075B-67	US-09-641-528-463
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27	30	31	33	3. 5. 5. 7.	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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Gaps
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            Sequence 9, Application US/60438805
GENERAL INFORMATION:
APPLICANT: KODADEK, THOMAS;
TITLE OF INVENTION: MIXED ELEMENT CAPTURE AGENTS;
FILE REFERENCE: UTSD:935USP1;
CURRENT APPLICATION NUMBER: US/60/438,805;
CURRENT FILNG DATE: 2003-01-09;
NUMBER OF SEQ ID NOS: 12;
SOFTWARE: Patentin Ver. 2.1;
SEQ ID NO 9
                                                                                                                                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
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US-60-438-805-9
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1 PLSQETFSDLWKLLPEN 17

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| Sequence 141, Application US/10211088 |
| APPLICANT: Bright, Gary R. |
| APPLICANT: Premkumar, D. David |
| APPLICANT: Chan, Yih-Tai |
| TITLE OF INVENTION: Novel Fusion Proteins And Assays For Molecular Binding |
| FILE REFERENCE: 01-102-US |
| CURRENT FAPLICATION NUMBER: 02/002-015 |
| PRIOR APPLICATION NUMBER: 00/309, 395 |
| PRIOR FILING DATE: 2001-08-01 |
| PRIOR PELING DATE: 2001-08-01 |
| PRIOR FILING DATE: 2001-12-13 |
| NUMBER OF SEQ ID NOS: 366 |
| SEQ ID NO 141 |
| SEQ ID NO 141 |
| LENGTH: 15 |
| LENGTH: 15

TYPE: PRT

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Gaps
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GENERAL INFORMATION:
APPLICANT: Charles, Ian G.
APPLICANT: Xu, Weiming
APPLICANT: Liu, Lizhi
TITLE OF INVENTION: Unducible Screen for Drug Discovery
TITLE OF INVENTION: Unducible Screen for Drug Discovery
FILE REFERENCE: HO-PO2380US0
CURRENT FILING DATE: 2000-07-28
FRIOR APPLICATION NUMBER: US/10/049,428
PRIOR FILING DATE: 1999-07-30
PRIOR PLING DATE: 1999-07-30
PRIOR PLING DATE: 1200-06-30
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                APPLICANT: Sidney, John
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Southwood, Scott
TITLE OF INVENTION: HLA Binding Peptides and Their Uses
FILE REPREBRUE: 3953-20019.20
CURRENT APPLICATION NUMBER: US/9/9/189,702A
CURRENT APPLICATION NUMBER: US 08/205,713
FRIOR FILING DATE: 1994-03-04
NUMBER OF SEQ ID NOS: 380
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 56.2%; Score 59; DB 5; Length 11; Best Local Similarity 100.0%; Pred. No. 0.0046; Matches 11; Conservative 0; Mismatches 0; Indels
Indels
0;
  0; Mismatches
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                                                                                                                                                                                                                                 ; Sequence 286, Application US/09189702A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                          APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
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Matches 9, Conservative
Matches 13; Conservative
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                                                        4 SQETFSDLWKLLP 16
                                                                                                     1 SOETFSDLWKLLP 13
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US-09-189-702A-286
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US-10-049-428-6
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GENERAL INFORMATION:
APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION:
TITLE OF INVENTION: PRANSCRIPTIONAL REGULATION
FILE REFERENCE: N73477C GCW
CURRENT APPLICATION WUMBER: US/09/701,080C
CURRENT APPLICATION NUMBER: GB 9811303.8
PRIOR APPLICATION NUMBER: GB 9811303.8
PRIOR APPLICATION NUMBER: GB 9900157.0
SROFTWARE: PATENT APPLICATION NUMBER: GB 9800157.0
FRIOR APPLICATION NUMBER: GB 9900157.0
SROFTWARE: PATENT APPLICATION NUMBER: GB 9800157.0
FRIOR APPLICATION NUMBER: GB 9900157.0
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GENERAL INFORMATION:
APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT APLICATION NUMBER: US 9811303.8
PRIOR APPLICATION NUMBER: GB 9811303.8
PRIOR FILING DATE: 1998-05-26
PRIOR FILING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Description of Artificial Sequence: derived from p53
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66.7%; Score 70; DB 5; Length 13;

Best Local Similarity 100.0%; Pred. No. 9.6e-05;

Matches 13; Conservative 0; Mismatches 0; Indels
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                                                                                                                                    77.1%; Score 81; DB 6; Length 15; 100.0%; Pred. No. 2e-06; tive 0; Mismatches 0; Indels
                                                  , OTHER INFORMATION: Binding domain US-10-211-088-141
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           ORGANISM: Artificial sequence
                                                                                                                                    Query Match
Best Local Similarity 100.0
Matches 15; Conservative
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US-09-701-080C-22
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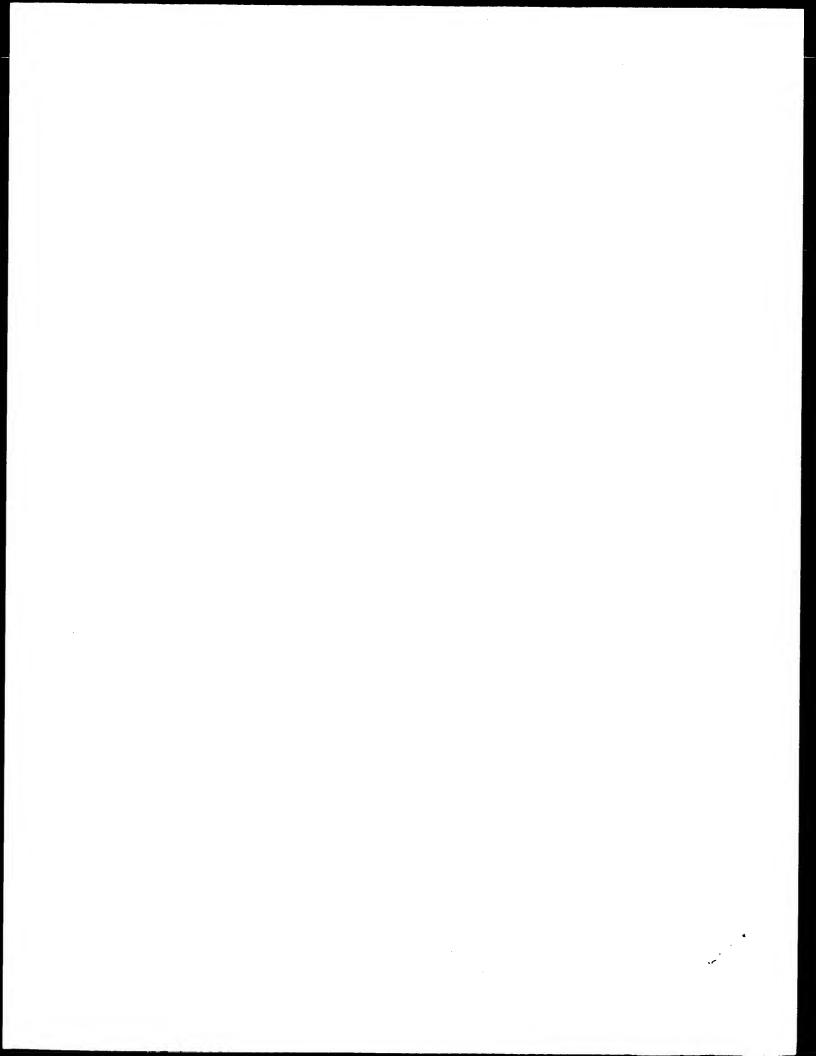
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TITLE OF INVENTION: IMMUNE- AND COLON- RELATED TISSUES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity 77.8
Matches 7; Conservative
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ORGANISM: Homo sapiens
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Best Local Similarity
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5 ESFSDELWK 13
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                                                                                                                                                                                                                                                                                                                                                                                                         SEQ ID NO 53
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TITLE OF INVENTION: NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPREMY3, EXPRESSED HIGHI
                                                                                          APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 I
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILLE REPERENCE: N134770 GCW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT FILLING DATE: 1098-05-26
PRIOR FILLING DATE: 1998-05-26
PRIOR FILLING DATE: 1998-01-05
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PARCENT NOS: 36
SOFTWARE: PARCENT NOS: 36
SEQ ID NO: 36
SEQ ID NO: 36
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; OTHER INFORMATION: Description of Artificial Sequence:derived from p53
US-09-701-080C-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        | Sequence 7. Application US/10049428 |
| GENERAL INFORMATION: | APPLICANT: Charles, Ian G. |
| APPLICANT: Charles, Ian G. |
| APPLICANT: Charles, Ian G. |
| APPLICANT: Liu, Lizhi |
| ITLE OF INVENTION: Unducible Screen for Drug Discovery |
| ITLE OF INVENTION: Unducible Screen for Drug Discovery |
| FILE REFERENCE: HO-P02380US0 |
| CURRENT APPLICATION NUMBER: US/10/049,428 |
| CURRENT APPLICATION NUMBER: GB 9918077 |
| PRIOR FILING DATE: 1999-07-30 |
| PRIOR FILING DATE: 1999-07-30 |
| PRIOR FILING DATE: 2000-06-30 |
| NUMBER OF SEQ ID NOS: 7 |
| SOFTWARE: PatentIn Version 3.1 |
| LENGTH: 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   38.1%; Score 40; DB 5; Length 7; 100.0%; Pred. No. 3.5e+05; tive 0; Mismatches 0; Indels
                                              Sequence 12, Application US/09701080C GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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Matches 7; Conservative
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RESULT 7 /
US-09-701-080C-12
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APPLICANT: LifeSpan Biosciences
APPLICANT: LifeSpan Biosciences
APPLICANT: LifeSpan Biosciences
APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Glenna C.
APPLICANT: Rough, Christine L.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 1920-4-4
CURRENT APPLICATION NUMBER: 2010-12-19
PRIOR PILING DATE: 2000-12-19
PRIOR FILING DATE: 2000-12-19
SPRIOR FILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 2292
SOFTWARE: Patentin version 3.1
SEQ ID NO 1177
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APPLICANT: Kast, W. Martin
APPLICANT: Southwood, Scott
APPLICANT: Epimmune, Inc.
TITLE OF INVENTION: HLA Binding Peptides and Their Uses
FILE REFERENCE: 39963-20019.20
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53.3%; Pred. No. 1.6e+02;
tive 2; Mismatches 2;
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CURRENT APPLICATION NUMBER: US/10/268,332
CURRENT APPLICATION NUMBER: US/10/268,332
CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: US. 60/235,713
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US. 60/261,783
PRIOR PLING DATE: 2001-01-16
PRIOR PLING DATE: 2001-01-16
PRIOR PLING DATE: 2001-01-13
PRIOR PELING DATE: 2001-01-13
PRIOR PELING DATE: 2001-08-17
PRIOR APPLICATION NUMBER: U.S. 60/313,171
PRIOR APPLICATION NUMBER: U.S. 60/313,171
PRIOR PELING DATE: 2001-09-17
PRIOR PLING DATE: 2001-09-26
NUMBER OF SEQ ID NOS: 66
SOFTWARE: PatentIn version 3.1
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APPLICANT: Darimont, Descrite
APPLICANT: Peng, Weijun
APPLICANT: Feng, Weijun
APPLICANT: Feng, Weijun
APPLICANT: Feng, Weijun
APPLICANT: Rushner, Peter
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Nachard
APPLICANT: Wagner, Nachard
APPLICANTION: OOACTIVACR BINDING
FILE REFERENCE: 9811-008-999;
CURRENT APPLICATION NUMBER: US 60/079,956
PRIOR APPLICATION NUMBER: US 60/079,956
PRIOR PILING DATE: 1998-03-30
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.0
SEQ ID NO 31
LENGTH: 12
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                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
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                                                                                                                                                                                                                                                                ; OTHER INFORMATION: p53.24V9 peptide 1323.02
US-09-189-702A-7
CURRENT APPLICATION NUMBER: US/09/189,702A
CURRENT FILING DATE: 1998-11-10
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1994-03-04
NUMBER OF SEQ ID NOS: 380
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
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OTHER INFORMATION: Leu --> Arg (L454R)
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OTHER INFORMATION: Leu --> Arg (L456R)
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; LOCATION: (8)..(8);
; OTHER INFORMATION: Glu -->Lys (E457K)
US-09-281-117A-31
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GENERAL INFORMATION:
APPLICANT: Baxter, John
                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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Best Local Similarity 60.0%
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LOCATION: (7)..(7)
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US-09-281-717A-33
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APPLICANT: Darimont, Beatrice
APPLICANT: Fletterick, Robert
APPLICANT: Fletterick, Robert
APPLICANT: Fletterick, Robert
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Brian
                                                                              APPLICANT: Darimont, Beatrice
APPLICANT: Darimont, Beatrice
APPLICANT: Darimont, Beatrice
APPLICANT: Feng, Weijun
APPLICANT: Feng, Weijun
APPLICANT: Feteraick, Robert
APPLICANT: West, Brian
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Reith
ITLE OF INVENTION: COACTIVAOR BINDING
ITLE OF INVENTION: COACTIVAOR BINDING
ITLE OF INVENTION: UNMBER: US/09/281,717A
CURRENT APPLICATION NUMBER: US 60/079,956
PRIOR FILING DATE: 1998-03-30
NUMBER OF SEQ ID NOS: 60
SOFTWARE PATENTIN VERSION 3.0
SEQ ID NO S: 60
SEQ ID NO S: 61
LENGTH: 12
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29.5%; Score 31; DB 5; Length 12;
Best Local Similarity 60.0%; Pred. No. 1.4e+02;
Matches 6; Conservative 0; Mismatches 4; Indels
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Sequence 33, Application US/09281717A GENERAL INFORMATION:
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Matches 6; Conserv
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### SPEICANT: Shidaw, Andrew
### APPLICANT: Shidaw, Andrew
### APPLICANT: Shidaw, Andrew
### APPLICANT: Shidaw Confered
### APPLICANT: Creene, Geoffrey
### APPLICANT: Greene, Geoffrey
### APPLICANT: Greene, Geoffrey
### TITLE OF INVENTION: ACTIVITY
### TITLE OF INVENTION: ACTIVITY
### TITLE OF INVENTION: ACTIVITY
### FILE REPERBNCE: 9811-013-99
### CURRENT PILING DATE: 1999-10-01-39
### PRIOR PILING DATE: 1999-10-30
### PRIOR PILING DATE: 1999-10-30
### PRIOR PILING DATE: 1999-12-16
### PRIOR PILING DATE: 1999-13-10
### PRIOR PILING DATE: 1990-13-10
### PRIOR PILING DATE: 1990-13-10
### PRIOR PILING DATE: 1990-13-13-10
### PRIOR PILING DATE: 1990-13-10
### PRIOR PILING DATE: 19
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74, Appl
8, Appli
10, Appl
11, Appl
16, Appl
                                                                                                                            February 12, 2003, 11:13:52; Search time 377.625 Seconds (without alignments) 32.439 Million cell updates/sec
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1: \cgn2 6/ptodata/1/paa/PCTUS COMB.pep:*

2: \cgn2 6/ptodata/1/paa/PCTUS COMB.pep:*

3: \cgn2 6/ptodata/1/paa/USO8 COMB.pep:*

4: \cgn2 6/ptodata/1/paa/USO8 COMB.pep:*

5: \cgn2 6/ptodata/1/paa/USO81 COMB.pep:*

7: \cgn2 6/ptodata/1/paa/USO81 COMB.pep:*

7: \cgn2 6/ptodata/1/paa/USO82 COMB.pep:*

10: \cgn2 6/ptodata/1/paa/USO83 COMB.pep:*

11: \cgn2 6/ptodata/1/paa/USO84 COMB.pep:*

12: \cgn2 6/ptodata/1/paa/USO85 COMB.pep:*

11: \cgn2 6/ptodata/1/paa/USO85 COMB.pep:*

12: \cgn2 6/ptodata/1/paa/USO87 COMB.pep:*

13: \cgn2 6/ptodata/1/paa/USO87 COMB.pep:*

14: \cgn2 6/ptodata/1/paa/USO87 COMB.pep:*

15: \cgn2 6/ptodata/1/paa/USO92 COMB.pep:*

16: \cgn2 6/ptodata/1/paa/USO93 COMB.pep:*

17: \cgn2 6/ptodata/1/paa/USO93 COMB.pep:*

18: \cgn2 6/ptodata/1/paa/USO93 COMB.pep:*

10: \cgn2 6/ptodata/1/paa/USO93 COMB.pep:*

10: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

10: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

20: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

21: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

22: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

23: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

24: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

25: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

26: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

27: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

28: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

29: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

20: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

21: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

22: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

23: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

24: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

25: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

26: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

27: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*

28: \cgn2 6/ptodata/1/paa/USO95 COMB.pep:*
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/cgn2_6/ptodata/1/paa/US60_COMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-214-371-74
US-10-155-059-8
US-10-155-059-10
US-10-155-059-11
US-10-155-059-11
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                                                                                                                                                                                                                                                                                                                                                                           4569144 seqs, 644733110 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                  OM protein - protein search, using sw model
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1 PPLSQETFSDLWKLLPENG 19
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Match Length
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Maximum DB seq length: 25
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Sequence 1, Appli
Sequence 3, Appli
Sequence 1109, Ap
Sequence 1109, Ap
Sequence 1109, Ap
Sequence 1469, Ap
Sequence 2, Appli
Sequence 3, Appli
Sequence 5, Appli
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Sequence 13, Appl
Sequence 26, Appl
Sequence 4, Appli
Sequence 4, Appli
Sequence 14, Appli
Sequence 18, Appli
Sequence 6, Appli
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                                   Sequence 14, App
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Sequence 20, Appl
                                                                                                       Sequence 13,
                                                                                                                                Sequence 19,
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                                                     Sequence 15,
                                                                     Sequence 73
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Sequence 8,
                                                                                     Sequence 7
                                                                                                                       Sequence
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| GENERAL INFORMATION: | APPLICANT: Lane, David Philip |
| TITLE OF INVENTION: MATERIALS AND METHODS RELATING TO |
| TITLE OF INVENTION: MATERIALS AND METHODS RELATING TO |
| TITLE OF INVENTION: INHIBITING THE INTERACTION OF p53 AND MDM2 |
| FILE REFERENCE: MEWB25.001APC |
| CURRENT PELLING DATE: 1200-01-19 |
| PRIOR PAPLICATION NUMBER: PCT/GB98/01144 |
| PRIOR FILING DATE: 1998-04-20 |
| PRIOR FILING DATE: 1997-04-22 |
| NUMBER OF SEQ ID NOS: 12 |
| SOFTWARE: PSELSEQ for Windows Version 4.0 |
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US-09-458-297-1109
US-09-458-297-1469
US-09-458-297A-1109
US-09-9458-297A-1469
US-09-936-956-3
US-09-936-956-3
US-08-277-666-5
US-08-277-666-5
US-08-277-666-5
US-08-277-66-5
US-08-5-5-5-5
US-08-5-5-5-5-5
US-08-5-5
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100.0%; Pred. No. 7.8e-09;
ive 0; Mismatches 0;
                              US-10-155-059-14

US-10-155-059-15

US-09-214-371-73

US-09-214-371-1

US-10-155-059-13

US-10-155-059-13

US-08-277-660-1

US-08-277-660-1

US-08-244-273-3

US-08-424-273-3
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US-09-936-956-13
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US-08-398-139B-8
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US-08-132-284-4
US-08-398-139B-4
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                US-09-403-440A-2
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SEQ ID NO 2
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Sequence 8, Application US/10155059
GENERAL INPORMATION:
APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODIS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                    APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Bcheverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
PILE REFERENCE: 4-20937/A/PCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ő
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94.3%; Score 99; DB 16; Length 18; 100.0%; Pred. No. 6.3e-08; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible OPERATING SYSTEM: Windows SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-MAY-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
                                                                                                                                                                                                                                                              CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
; Sequence 74, Application US/09214371B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 PPLSQETFSDLWKLLPEN 18
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Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
                                                  Lane, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 02110
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LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                     APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
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APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
                                                                                                                                                                                                                                                                                                                                                                    0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94.3%; Score 99; DB 25; Length 20; 100.0%; Pred. No. 7.2e-08;
                                                                                                                                                                                                                                                                                                               94.3%; Score 99; DB 25; Length 20; 100.0%; Pred. No. 7.2e-08; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CURRENT ITEL COMPACTURE
COMPUTER: IBM Compactible
COPERATING SYSTEM: Windows
SOFTWARE: FastSEM: Windows
SOFTWARE: FastSEM: Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,10/155,059
FILING DATE: 24-349-2002
CLASSIPICATION: <Unknown>
PRIOR APPLICATION NUMBER: US,09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: G0/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                            TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 8:
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                   TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 10, Application US/10155059; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 20 amino acids
                                                                                                                                                                LENGTH: 20 amino acids
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INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single
                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 02110
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                   1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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NUMBER OF SEQUENCES: 27
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Best Local Similarity 100.0
Matches 18; Conservative
                                                                            CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 20005
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-10-155-059-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: V
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 7
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US-10-155-059-16
i Sequence 16, Application US/10155059
i GENERAL INFORMATION:
i APPLICANT: Kaelin, William
i APPLICANT: Kaelin, William
i TITLE OF INVENTION: METHODS OF TREATMENT USING
i TITLE OF INVENTION: MES-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                                                                                      RESULT 5
US-10-155-059-11
Sequence 11, Application US/10155059
GENERAL INFORMATION:
Habelicant Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: MBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ö
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       Gaps
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       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: FastSEQ for Windows Version 2.0b CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIORICATION DATE: US/10/155,059
FILING DATE: 24-May-2002
CLASSIPICATION: UNKNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 37400
       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS: ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible OPERATING SYSTEM: Windows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 20 amino acids
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 PPLSQETFSDLWKLLPEN 19
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                                                                 1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
       Conservative
                                                                                                                                                                                                                                                                                                                                                                              CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-155-059-11
       18;
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94.3%; Score 99; DB 25; Length 20; 100.0%; Pred. No. 7.2e-08; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                            COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
SOFTWARE: FastSEG for Windows Version 2.0b
CURRBY APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24 May-2002
CLASSIFICATION: <u href="checkbase">cunscientes</a>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REPERENCE/DOCKET NUMBER: 47400
TELECOMFUNICATION INFORMATION:
TELEPAK: 617-345-6054
TELEFAK: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 16:
                   ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 18, Application US/08244776; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Zentgraf, Hanswalter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                              MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 16:
                                                                                                                    COUNTRY: USA
ZIP: 02110
COMPUTER READABLE FORM:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PPLSQETFSDLWKLLPEN 18
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US-08-424-273-1

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TITLE OF INVENTION: P53 PROTEIN FRAGMENTS AND USE THEREOF FOR THILE OF INVENTION: DETECTING AND MONITORING DISEASED CONDITIONS NUMBER OF SEQUENCES: 15 CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              94.3%; Score 99; DB 6; Length 25; 100.0%; Pred. No. 9.3e-08; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Jacobson, Price, Holman & Stern, PLLC STREET: 400 Seventh Street, N.W.
                                                      PULICATION DATA
APPLICATION DATA
APPLICATION DATA
PFLICATION NUMBER: WO 94/08241
FILING DATE: 30-SEP-1993
APPLICATION NUMBER: DE P 42 32 823.3
FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: Ihnen, Jeffrey L.
REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 18644-94281-1
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READBLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Word Perfect 5.1 (ASC III)
CURRENT APPLICATION DATA:
APPLICATION WUMBER: US/08/424,273
FILING DATE: 28-APR-1995
CLASSIFCATION: 435
FILING DATE: 2-NOV-1992
FRIOR APPLICATION DATA:
APPLICATION NUMBER: FR 9213110
FILING DATE: 2-NOV-1992
FRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/FR 9301082
FILING DATE: 2-NOV-1993
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acide
APPLICATION NUMBER: US/08/244,476
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-424-273-1; Sequence 1, Application US/08424273; Sequence 1, Application APPLICANT: APPLICANT:
                                                                                                                                                                                                                                                                                                         TELEPHONE: 202-962-4810
TELEPAX: 202-962-8300
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
              FILING DATE: 15-SEP-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 PPLSQETFSDLWKLLPEN 18
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Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS:
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USA
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COUNTRY:
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TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
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      94.3%; Score 99; DB 8; Length 25; 100.0%; Pred. No. 9.3e-08; rive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMBUTE: IBM Compatible
COMBUTE: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows
SOFTWARE: FastSEQ for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION AURHOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STRANDENESS: single
; TOPOLOGY: linear
; SEQUENCE DESCRIPTION: SEQ ID NO: 14:
US-10-155-059-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                   RESULT

; Sequence 14, Application US/10155059

; GENERAL INFORMATION:

; APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 10
US-10-155-059-15; Sequence 15, Application US/10155059
; GENERAL INFORMATION:
. APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: MA
COUNTRY: USA
ZIP: 0210
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 27
                                                                                       1 PPLSQETFSDLWKLLPEN 18
                                                                                                                 2 PPLSQETFSDLWKLLPEN 19
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Best Local Similarity 94.4'
Matches 17, Conservative
                                             Conservative
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Query Match
Best Local Similarity
Matches 18; Conserv
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RESULT 13
US-10-155-059-13
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US-09-732-384-7
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          LENGTH: 22
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                                                                             FEATURE:
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JOST, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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GENERAL INFORMATION:
APPLICANT: Lane, David
APPLICANT: Lane, Dattger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Picksley, Stephen
APPLICANT: Picksley, Stephen
APPLICANT: Picksley, Stephen
APPLICANT: Picksppel, Hainz-Kurt
APPLICANT: Hochkeppel, Hainz-Kurt
APPLICANT: Hochkeppel, Hainz-Kurt
APPLICANT: Hochkeppel, Hainz-Kurt
APPLICANT: Hochkeppel, Hainz-Kurt
APPLICANT: Hordkeppel, Hainz-Kurt
APPLICANT: Hordkeppel, Hainz-Kurt
APPLICANT: Hure, Pascal
ITILE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REFERENCE: 4-20337/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR FILING DATE: 1999-03-26
PRIOR FILING DATE: 1999-03-26
NUMBER OF SEQ ID NOS: 83
SOFTWARR: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGRYT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELEPHONE: 617-345-6054
TELEPHONE: 617-345-6054
TELEPHONE: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                    NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                         ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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nes 17; Conservative
                                                                                                                                                                                                                                                COUNTRY: USA
                                                                                                                                                                                              CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-214-371-73
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NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE ANTIBODIES
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0
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                                                                    OTHER INFORMATION: Description of Artificial Sequence:peptide
                                                                                                                                                                                                     88.6%; Score 93; DB 16; Length 22; 100.0%; Pred. No. 6.9e-07; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87.6%; Score 92; DB 21; Length 19; 100.0%; Pred. No. 8.4e-07;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
OPERATING SYSTEM: Windows
OPERATING SYSTEM: Windows
OFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Yunn, Zhi-Min
INTILE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APLICATION VUMBER: U5/09/732,384
CURRENT APPLICATION NUMBER: 06/169,816
FRICH APPLICATION NUMBER: 60/169,816
FRICH FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PATENTING VET: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kaelin, William Jost, Christine TITLE OF INVENTION: METHODS OF TREATMENT USING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 13, Application US/10155059 GENERAL INFORMATION:
                                                                                     NAME/KEY: VARIANT
LOCATION: (1)
SOTHER INFORMATION: X = Biotin-Ser
US-09-214-371-73
TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 02110
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 PLSQETFSDLWKLLPEN 17
                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPE 17
                                                                                                                                                                                                                                                                                                                                        6 PPLSQETFSDLWKLLPE 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 PLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                     Query Match 88.6
Best Local Similarity 100.
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: MA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Furet, Pascal TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2 FILE REFERENCE: 4-2093/APCT CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION PRIOR PRIOR PLICATION PRIOR PLICATION PRIOR PLICATION NUMBER: PCT/EP97/03549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ó
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 85.7%; Score 90; DB 25; Length 19; Best Local Similarity 88.9%; Pred. No. 1.7e-06; Matches 16; Conservative 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               84.8%; Score 89; DB 16; Length 19; 94.1%; Pred. No. 2.5e.06; tive 1; Mismatches 0; Indels
                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <UNKnown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 13: US-10-155-059-13
                                                                                                                                                                                                   NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFRENSE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Lane, David
Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
Hochkeppel, Hehnz-Kurt
Garcia-Echeverria, Carlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/09214371B GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                TELEX: <Unknown>
MATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 94.1%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 PPLSQETFSELWNLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PLSQOTFSDLWKLLPEN 17
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US-09-214-371-1
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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SEQ ID NO 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT:
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RESULT 15

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                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OOPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
PPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Search completed: February 12, 2003, 11:24:05 Job time : 378.625 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Eisenstein, Rohald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                            ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                 ; Sequence 19, Application US/10155059; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 20 amino acids
                                                            APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 PLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 94.1
Matches 16; Conservative
                                                                                                                                                                    NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                     CITY: Boston
                                                                                                                                                                                                                                                                                             COUNTRY: USA
                                                                                                                                                                                                                                                                                                                    ZIP: 02110
                                                                                                                                                                                                                                                                           STATE: MA
US-10-155-059-19
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GenCore version 5.1.3
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OM protein - protein search, using sw model

Pebruary 12, 2003, 11:15:02 ; Search time 12.6667 Seconds
 (without alignments)
 38.323 Million cell updates/sec Run on:

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19 Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

140259 segs, 25548876 residues

42882 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/USOB_NEW_PUB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/PUT.NEW_PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
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8: /cgn2_6/ptodata/1/pubpaa/USO9_NEW_PUB.pep:*
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14: /cgn2_6/ptodata/1/pubpaa/USO0_NEW_PUB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

	Description	Semience 74. Appl	α	0	Sequence 11 Appl	Segmence 16 Appl	Sequence 14. Appl	2	, 5	, ,	~	, ,	16	Sequence 6, Appli	23	20.	181	12		¢ .	sednence 77, Appl
	ΙD	US-09-214-371-74	US-10-155-059-8	US-10-155-059-10	US-10-155-059-11	US-10-155-059-16	US-10-155-059-14	US-10-155-059-15	US-09-214-371-73	US-09-732-384-7	US-10-155-059-13	US-09-214-371-1	US-10-155-059-19	US-09-732-384-6	US-10-155-059-23	US-10-155-059-20	US-10-155-059-18	US-09-214-371-17	3-69-950-8D	115-10-155-059-22	77-600-661-01-60
	DB	10	12	12	12	12	12	12	10	10	12	10	12	10	12	12	12	10	10	12	1
	Query Match Length DB	18	20	20	20	20	20	20	22	19	19	19	20	15	16	20	20	12	15	18	1
ф	Query Match	94.3	94.3	94.3	94.3	94.3	89.5	89.5	98.6	87.6	85.7	84.8	81.9	73.3	70.5	70.5	68.6	62.9	61.9	57.1	:
	Score	66	66	66	66	66	94	94	93	92	90	89	98	77	74	74	72	99	65	9	,
	Result No.	H	7	e	4	S	9	7	6 0	σ	10	11	12	13	14	15	16	17	18	19	ì

Segmence 39. Appl	Seguence 7, Appli		Sequence 24. Appl	41	40,	Sequence 25, Appl	26.	Sequence 33, Appl	34	17	m	Sequence 21, Appl	20	24	83	4	٦,	37,	38,	35,	Sequence 36, Appl	2. A			13, 7
10 US-09-214-371-39	9 US-10-024-123-7	10 US-09-214-371-7	10 US-09-214-371-24	10 US-09-214-371-41	10 US-09-214-371-40	10 US-09-214-371-25	10 US-09-214-371-26	10 US-09-214-371-33	10 US-09-214-371-34	12 US-10-155-059-17	10 US-09-019-679-3	12 US-10-155-059-21	10 US-09-214-371-20	12 US-10-155-059-24	10 US-09-214-371-83	10 US-09-732-384-4	12 US-10-155-059-1	10 US-09-214-371-37	10 US-09-214-371-38	10 US-09-214-371-35	10 US-09-214-371-36	12 US-10-155-059-2	9 US-09-840-277-66	10 US-09-920-552-110	10 US-09-214-371-13
16	14	12	12	16	16	12	12	14	14	10	16	11	12	15	9	v	9	6	0	10	10	9	12	17	œ
9 56.2	7 54.3	5 52.4	4 51.4	4 51.4	3 50.5	9 46.7	9 46.7	8 45.7	8 45.7	7 44.8	7 44.8	4 41.9	4 41.9	2 40.0	6 34.3	6 34.3	6 34.3	6 34.3	6 34.3			2 30.5	2 30.5	m	1 29.5
2	S	S	S	ß	2	4	4	4	4	4	4	4	4	4	e	m	٣	m	3	m	m	m	m	'n	m
20	21	22	23	24	25	56	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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Gaps
                                                                                                                                                                                                                                        APPLICANT: Hochkepel, Heinz-Kurt
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REFERENCE: 4-20937/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 94.3%; Score 99; DB 10; Length 18; Best Local Similarity 100.0%; Pred. No. 5.5e-09; Matches 18; Conservative 0; Mismatches 0; Indels
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                         ; Sequence 74, Application US/09214371B
; Patent No. US20010018511A1
; GENERAL INFORMATION:
; APPLICANT: Lane, David
; APPLICANT: Bottger, Volker
; APPLICANT: Bottger, Angelica
; APPLICANT: Picksley, Stephen
; APPLICANT: Picksley, Stephen
; APPLICANT: Garcia-Echeverria, Carlos
; APPLICANT: Garcia-Echeverria, Carlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 74
LENGTH: 18
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US-09-214-371-74
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1 PPLSQETFSDLWKLLPEN 18 RESULT 2 US-10-155-059-8 qq

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; Sequence 8, Application US/10155059 ; Patent No. US20020147173A1

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Jost, Christine
TITLE OF INVENTION: METHODISS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
               APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODIES OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FeatsEng for Windows Version 2.0b
SURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 47400
TELEPHONE: 1NFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
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SEQUENCE DESCRIPTION: SEQ ID NO: 8:
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
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Patent No. US20020147173A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 8: SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
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                                                                                                                                         NUMBER OF SEQUENCES:
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                                                                                                                                                                                                                                          CITY: Boston
                                                                                                                                                                                                                                                                                        COUNTRY: USA
GENERAL INFORMATION:
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APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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94.3%; Score 99; DB 12; Length 20;
Best Local Similarity 100.0%; Pred. No. 6.2e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
               COMPUTER: IBM Compatible
COMPRING SYSTEM: Windows
SOFTWARE: FastSENG for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24 May-2002
CLASSIRICATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1999
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
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COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
ODERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                      NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPRAK: 617-345-6054
TELEFAX: 617-345-1300
TELEX: CURROWN:
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SEQUENCE DESCRIPTION: SEQ ID NO: 10:
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CORRESPONDENCE ADDRESS:
ADDRESSE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Eisenstein, Ronald I
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; Sequence 11, Application US/10155059
; Patent No. US20020147173A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
ZIP: 02110
COMPUTER READABLE FORM:
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ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
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         94.3%; Score 99; DB 12; Length 20; 100.0%; Pred. No. 6.2e-09; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM: Windows
SOFTWARE: FRISHED for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
                                                                                                                                                                                                                                                                                                             APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
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ADDRESSEE: Nixon Peabody LLP
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CITY: Boston
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CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                               ; Sequence 14, Application US/10155059; Patent No. US20020147173A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: <UNKNOWN>
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 27
                                                                                               1 PPLSQETFSDLWKLLPEN 18
                                                                                                                       2 PPLSQETFSDLWKLLPEN 19
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                                                    18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
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Best Local Similarity
Matches 17; Conserva
                               Best Local Similarity
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US-10-155-059-14
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           Query Match
                                                    Matches
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JOSt, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NIS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: 24-May-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
ATTORNEY/AGENT INFORMATION:
NAME: Bisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                        INFORMATION FOR ESQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 11:
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 16:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20 amino acids
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INFORMATION FOR SEQ ID NO: 16:
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                2 PPLSQETFSDLWKLLPEN 19
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
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ZIP: 02110
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US-10-155-059-16
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; Sequence 13, Application US/10155059; Patent No. US20020147173A1; GENERAL INFORMATION:
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US-09-214-371-73
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
SEQ ID NO 73
LENGTH: 22
TYPE: PT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: USA
ZIP: 02110
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 PLSQETFSDLWKLLPEN 17
                                                                                                                                                                                                                                                                                                      1 PPLSQETFSDLWKLLPE 17
                                                                                                                                                                                                                                                                                                                                  6 PPLSQETFSDLWKLLPE 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-10-155-059-13
                                                                                                                                             LOCATION: (1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
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                                          NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
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94.4%; Pred. No. 3.5e-08;
tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: CURROWN>
PRIOR APPLICATION DATA:
      JOST, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/09/081,975
PILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Bisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 15: US-10-155-059-15
                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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Garcia-Echeverria, Carlos
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                                                                  ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                        MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS: single
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Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
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Best Local Similarity 94.44
Matches 17; Conservative
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TITLE OF INVENTION: Inhib
FILE REFERENCE: 4-20937/A
                                                                                     NUMBER OF SEQUENCES:
                                                                                                                                                                     CITY: Boston
                                                                                                                                                                                                              COUNTRY: USA
                                                                                                                                                                                                                                 ZIP: 02110
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APPLICANT:
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APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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FEATURE: OTHER INFORMATION: Description of Artificial Sequence:peptide NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Ouery Match 87.6%; Score 92; DB 10; Length 19; Best Local Similarity 100.0%; Pred. No. 6.6e-08; Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                            Query Match

88.6%; Score 93; DB 10; Length 22;
Best Local Similarity 100.0%; Pred. No. 5.5e-08;
Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 7, Application US/09732384

Patent No. US/00201329771

GENERAL INCRATION:

APPLICANT: Yuan, Zhi-Min

APPLICANT: Yuan, Zhi-Min

TITLE OF INVENTION: Inhibition of p53 Degradation

FILE REFERENCE: 21508-044

CURRENT APPLICATION UNMER: US/09/732,384

CURRENT APPLICATION NUMBER: 60/169,816

PRIOR APPLICATION NUMBER: 60/169,816

PRIOR FILING DATE: 199912-08

NUMBER OF SEQ ID NOS: 10

SOFTWARE: PatentIn Ver. 2.1
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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1 PLSQQTFSDLWKLLPEN 17
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ZIP: 02110
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Best Local Similarity
                                                                         RESULT 12
US-10-155-059-19
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US-09-732-384-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Picksley, Stephen APPLICANT: Picksley, Stephen APPLICANT: Chene, Patrick APPLICANT: Chene, Patrick APPLICANT: Hochkeppel, Heina, Carlos APPLICANT: Earcia-Echeverria, Carlos APPLICANT: Furet, Pascal TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2 FILE REPERENCE: 4-2093/A/PCT CURRENT APPLICATION NUMBER: US/09/214,371B CURRENT FILING DATE: 1999-03-26 PRIOR FILING DATE: 1999-03-36 PRIOR FILING DATE: 1997-07-04 NUMBER OF SEQ ID NOS: 83
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            85.7%; Score 90; DB 12; Length 19; 88.9%; Pred. No. 1.3e-07;
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FASTSED for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION AURIEN: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/46,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                NAME: Elsentein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Picksley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Garcia-Echeverria, Carlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 1, Application US/09214371B; Patent No. US20010018511A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 19 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
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ORGANISM: Artificial Sequence
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SEQ ID NO 1
LENGTH: 19
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Matches 16; Conservative
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US-09-214-371-1
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NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OOFERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                         JOST, Christine TITLE OF INVENTION: METHODS OF TREATMENT USING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 6, Application US/09732384
Patent No. US/20020132977A1
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, JiJie
TITLE OF INVENTION: Inhibition of p53 Degradation
TITLE OF INVENTION: Inhibition of p53 Degradation
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: CURKNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE DESCRIPTION: SEQ ID NO: 19:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                                                                                                                      STREET: 101 Federal Street
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TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
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                                                                                                                                                                                     ANTIBODIES
; Sequence 19, Application US/10155059; Patent No. US20020147173A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                 APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single
                                                                                                                                                                                                             NUMBER OF SEQUENCES: 27
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PRIOR FILING DATE: 1999-12-08

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APPLICANT: Kaelin, William
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TELEX: <Unknown>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS:
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Best Local Similarity 93.3°
Matches 14; Conservative
         2 PLSQETFSDLWKLLPE 17
                                               1 PLSOETFEDLWSLLPD 16
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                                                                                                                                    US-10-155-059-20
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                                                                                                                 RESULT 15
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Jost, Christine
JITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
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                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: Protein COTHER INFORMATION: fragment not in inhibitory p53 polypeptide US-09-732-384-6
                                                                                                                                                                                                                                   Ouery Match 73.3%; Score 77; DB 10; Length 15; Best Local Similarity 100.0%; Pred. No. 9.4e-06; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 16;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-MAY-2002
CLASSIFICATION - CURROWN>
PRIOR SPELICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 23:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Eisenstein, Ronald I
REGIGTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 23, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 16 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Disket
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEOUENCES:
                                                                                                                                                                                                                                                                                                                          5 QETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                       1 QETFSDLWKLLPEN 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: USA
                      SEQ ID NO 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-155-059-23
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TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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                                                                                                                                                                                                                                                                                                                                                                                                                            MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEG for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Search completed: February 12, 2003, 11:25:07 Job time : 12.6667 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 20:
                                                                                                                                                                                                   NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
Sequence 20, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 11:10:52; Search time 47.5 Seconds (without alignments) 53.300 Million cell updates/sec Run on:

Title: Perfect score:

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19

Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

908470 segs, 133250620 residues Searched:

320064 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Databası

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	p53 homologue TIP p53 homologue TIP p53 N-terminal pep Immunodominant epi Human p53 amino ac p53/RB interaction p53 biotinylated N Human p53 peptide Immunodominant epi Immunodominant epi
SUMMARIES ID	AAW82321 AAW32319 AAW3228 AAR31907 AAR51879 AAW47074 AAW3727 AAW3727 AAR5409
% Query Match Length DB	250 118 118 118 118 118 118 118 118 118 11
% Query Match L	100.0 100.0 94.3 94.3 92.4 92.4 77.1 73.3
Score	1005 1005 999 97 93 77 77
Result No.	10 10 10

Disclosure, Fig 1, 52pp, English.

p53 protein bindin Peptide #1: Unide	3a mot	inding	S	13. U	Peptide #7. Unide	lated	#11.	#4.		II SS	TRAM-interaction m	Peptide #6. Unide	#	#1	e #2.	ın	Human oncogenic pr	⊢	Mdm/hdm antagonist	Mdm/hdm antagonist	Mdm/hdm antagonist	Human oncogenic pr		Tumour suppressor		p53 DR supermotif		Human oncogenic pr		Mdm/hdm antagonist	Mdm/hdm antagonist	Mdm/hdm antagonist	Human oncogenic pr
AAB29157	AAG89500	AAG89730	AAY45227	AAB29159	AAB29163	ABB05528	AAB29167	AAB29160	AAB29161	AAY99001	AAY57799	AAB29162	AAB29164	AAB29149	AAB29158	AAW37181	AAW37188	AAB17075	AAB17087	ABB73170	ABB73182	AAW37201	AAR83012	AAY95859	ABB09246	AAG89447	AAM52269	AAW37190	AAB17077	AAB17089	ABB73172	ABB73184	AAW37203
17	22	22	20	21	21	23	21	21	21	21	21	21	21	21		19					23		16	21	23	22	23		21	21	23		13
15	15	15	17	15	15	14	15	15	15	15	13	15	12	16	16	12	12	12	15	12	12	16	12	15	15	15	11	12	12			12	
77 73.3	73	77 73.3	73.	70.	70.	69	9	67.	71 67.6	67.	.99	65.	65 65.	2	.5 63.	9	9	9	6 62.	9		62.	5 61.	5 61.	9	2 59.	1 5	1 58.	1 58.	1 58.	1 5	1 5	61 58.1
11	13	14	15	16	17	18	19	20	21	22	23	24	25	56	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.
                                                                                                                                                                                                                                           New substance with a mdm2 binding domain and coupling partner -useful for stabilising in cells without an efficient mdm2-mediated degradation pathway
              AAW82321 standard; Peptide; 19 AA.
                                                                                                                                                       98WO-GB01140.
                                                                                                                                                                      97GB-0008089.
                                                (first entry)
                                                                  p53 homologue TIP peptide.
                                                                                                                                                                                                                          WPI; 1998-609975/51.
                                                                                                                                                                                         (UYDU-) UNIV DUNDEE
                                                                                                                    WO9847919-A1
                                                                                                                                                      20-APR-1998;
                                                                                                                                                                       22-APR-1997;
                                                 22-FEB-1999
                                                                                                                                     29-OCT-1998
                                                                                                   Synthetic.
                                AAW82321;
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     RESULT 1
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X0000000000000X8

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100.0%; Pred. No. 5.7e-10;

Best Local Similarity

Gaps

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This sequence is a peptide homologue of a region of p53 which binds to mana. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells on or overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.
This sequence is a peptide homologue of a region of p53 which binds to madm2. This peptide is used in the construction of a movel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells on or overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 tunction. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New agents which inhibit interaction of p53 and mdm2 - useful for activating p53, e.g. for treating cancers, viral conditions or other conditions associated with non functional p53 or mdm2
                                                                                                                                                                                                                                                                                                                        Length 19;
                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                   100.0%; Score 105; DB 19;
100.0%; Pred. No. 5.7e-10;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                            1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPENG 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  p53 homologue TIP peptide.
                                                                                                                                                                                                                                                                                                                                           Best Local Similarity 100. Matches 19; Conservative
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                                                                                                                                                                                                                                                                              19 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Synthetic.
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                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW82319;
                                                                                                                                                                                                                                                                                                                          Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lane DP;
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This represents a p53 N-terminal peptide fragment used in an Elisa TIP assay for analysing the interaction between human oncogenic protein MDM2 can dip53. The invention provides peptide derivatives capable of binding to the human MDM2. These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated can peptides may be used to identify molecules that bind compared to identify and design inhibitors of MDM2, p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia conteractions, especially tumours and viral infections. The peptides can interactions, especially tumours and viral infections. The peptides can be administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and caccumulation in normal cells. The peptides which mimic the MDM2 binding content in p53, have a significantly greater blocking activity compared content with wild-type p53.
                                                                                                                                                                                                                                                                                                           MDM2; oncogenic protein; p53; human; inhibition; interaction; cancer; tumour; diagnosis; binding; viral infection; Elisa TIP assay.
 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Compounds binding to MDM2 protein and inhibit its interaction with p53 - useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Boettger V, Chene P, Furet P, Garcia-echeverria C;
Lane D, Picksley S;
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 Indels
                                                                                                                                                                                                                                                                          p53 N-terminal peptide fragment for Elisa TIP assay.
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 0;
Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 34; 45pp; English.
                                                                                                                                                                 AAW37228 standard; peptide; 18 AA.
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                                      1 PPLSQETFSDLWKLLPENG 19
                                                                      1 PPLSQETFSDLWKLLPENG 19
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                                                                                                                                                                                                                                         20-JUL-1998 (first entry)
 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (NOVS ) NOVARTIS AG.
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Hochkeppel H,
                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 04-JUL-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       05-JUL-1996;
 19;
                                                                                                                                                                                                      AAW37228;
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 Matches
                                                                                                                                RESULT 3
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100.0%; Score 105; DB 19; Length 19;

Query Match

AAR5490 RESULT

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Zentgraf H;

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p53/RB interaction inhibiting peptide 6 (residues 11-29 of human p53).
                                                                                                                                                                                                                                                                                                                                                               Antibodies specific for p53 are detected by binding to immobilised fragments of the p53 gene product containing the antibody-binding region. Preferred fragments contain maino acids 1-241, 06-349, 40-393, 66-241, 66-393, 237-349, 237-393 and esp. 9-33, 37-52 or 368-386. See AAR51872-R51881 for sequences of these fragments.
                                                                                                                                                                                                                                        Non-radioactive detection of p53 specific antibodies - by capture on immobilised p53 or its fragments, then reaction with labelled second antibody, for diagnosis of tumours and suitable for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Identifying compounds that modulate interaction of p53 and Rb protein - or those that bind to Rb protein, used to induce apoptosis, specifically for treatment of tumours
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Retinoblastoma gene; RB; p53 protein; interaction; inhibitor; tumour; apoptosis; modulator; medicine; veterinary; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94.3%; Score 99; DB 15; I
100.0%; Pred. No. 7.1e-09;
ive 0; Mismatches 0;
                                                                                                                                                     Volkmann M,
                                                                                                                      (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM STIFTUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (CANC-) CANCER RES CAMPAIGN TECHNOLOGY
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                                                 93WO-EP02666
                                                                                   92DE-4232823
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 100.
Les 18; Conservative
                                                                                                                                                      Schranz P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1997-549887/50.
                                                                                                                                                                                         WPI; 1994-135732/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          25 AA;
                                                                                                                                                                                                           N-PSDB; AAQ62365
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens.
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                                                 30-SEP-1993;
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                14-APR-1994.
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                                                                                                                                                                                                                                                                                                   screening
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptides derived from the N-terminal (amino acids 1-112) or the C-terminal (amino acids 350-393) of protein p53 which specifically eact with anti-p53 antibodies in patients with cancer or precancerous conditions are claimed. The peptides (ARE4907-R54921) are useful for detecting and monitoring cancerous and precancerous
                                                                                                                                                                                                                        cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen; transformation-associated protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New immuno:dominant epitope(s) of protein p53 - for detecting and monitoring antibodies indicative of cancer and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human nuclear phosphoprotein p53; tumour suppressor gene product; anti-oncogene; cancer; tumour; antibody binding region; epitope.
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Pred. No. 7.1e-09;
0; Mismatches 0; Indels
                                                                                                                                                                                         Immunodominant epitope from p53 N-terminal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR51879 standard; Protein; 25 AA.
                                                                                 AAR54907 standard; peptide; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 4; Page 42; 62pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Soussi T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94.3%; S
100.0%;
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1 PPLSQETFSDLWKLLPEN 18
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (EURO-) LAB EUROBIO SA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lubin R,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1994-167463/20
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                                                                                                                      AAR54907;
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AAR51879

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Gaps

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Length 25; 0; Indels

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corresponds to residues 11-29 of human p53 (AAW47079). The interaction between p53 aprotein and a retinoblastoma (RB) protein. This peptide corresponds to residues 11-29 of human p53 (AAW47079). The interaction between p53 and Rb is found to be critical for determining whether or not a cell enters apoptosis. Apoptosis is prevented if interaction occurs. The interaction is between regions 1-71 or 290-393 of p53 and region 739-928 of Rb. The invention provides methods to identify compounds able to modulate interaction or binding between them These inhibitory and detecting interaction/binding between them These inhibitory compounds are used in human or veterinary medicine to modulate p53 activity and processes, specifically for inducing apoptosis in tumour cells (possibly also in cells infected by virus), in vivo or in vitro. Expression of these modulators by gene therapy methods is also contemplated. Other activities that can be affected are transcription repression, G1 arrest, DNA repair, homologous recombination and 23-5- exonuclease activity. Modulation of interaction with Rb may also
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               This peptide fragment of p53 is an inhibitor of the interaction between
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MDM2; oncogenic protein; p53; human; inhibition; interaction; cancer; tumour; diagnosis; binding; viral infection; Elisa P2 assay.
                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Compounds binding to MDM2 protein and inhibit its interaction with p53 - useful in, e.g. diagnosis and treatment of cancer and viral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Garcia-echeverria C;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           p53 biotinylated N-terminal peptide fragment for Elisa P2 assay.
                                                                                                                                                                                                                                                                                                                                                                                                 Score 97; DB 18; Length 19;
Pred. No. 1.1e-08;
1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Boettger V, Chene P,
Lane D, Picksley S;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Disclosure; Page 34; 45pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW37227 standard; peptide; 22 AA.
                                                                                                                                                                                                                                                                                                                                                                                                 92.4%;
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96GB-0014197.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 94.44
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (NOVS ) NOVARTIS AG.
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                                                                                                                                                                                                                                                                                                                                                            Sequence 19 AA;
                                                                                                                                                                                                                                                                                                                          stabilise p53.
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Hochkeppel H,
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05-JUL-1996;
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This p53 biotinylated N-terminal peptide fragment is used in an Elisa P2 assay for analysing the interaction between human oncogenic protein MDM2 and p53. The invention provides peptide derivatives capable of binding to the human MDM2. These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2 and to identify and design inhibitors of MDM2, p53 binding. They may also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia interactions, especially tumours and viral infections. The peptides can be administered masally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and site in p53, have a significantly greater blocking activity compared with wild-type p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New p53 regulatory protein (RB18A) useful as, e.g. sources of probes and primers to detect the transcription rate and abundance of RB18A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           wound healing; graft rejection; reperfusion injury;
myocardial infarction; stroke; traumatic brain injury;
meurodegenerative disease; ischaemia; toxaemia; infection; AIDS;
hepatitis; breast cancer; ovarian cancer; colon cancer; diagnosis;
therapy; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RB18A, p53 regulatory protein; apoptosis; neoplasia; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This synthetic peptide corresponds to amino acids 13-27 of human pags. It was used to raise anti-peptide p53.1 and anti-RB18A antibodies in rabbit. RB18A (see AAVG6309) is a novel p53 regulatory protein of the invention. The antibodies were used in
                                                                                                                                                                                                                                                                                                                                                                                               Length 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                             88.6%; Score 93; DB 19; L
100.0%; Pred. No. 5.7e-08;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human p53 peptide (aa13-27).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 PPLSQETFSDLWKLLPE 17
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                                                                                                                                                                                                                                                                                                                                                           22 AA;
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AAR54910 standard; peptide; 15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Peptides derived from the N-terminal (amino acids 1-112) or the C-terminal (amino acids 350-393) of protein p53 which specifically react with anti-p53 antibodies in patients with cancer or precancerous conditions are claimed. The peptides (AAR54907-R54921) are useful for detecting and monitoring cancerous and precancerous
                      and
   immunoscreenings that led to the isolation of a cDNA clone (see AAX59124) encoding human RB18A. The invention provides methods and compositions for the diagnostic and therapeutic applications of RB18A, in particular for the diagnosis, prevention or treatment of
                                                                                                                                                                                                                                                                                                                                                                                                                          cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New immuno:dominant epitope(s) of protein p53 - for detecting and monitoring antibodies indicative of cancer and precancerous
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                                                                                                                           77.1%; Score 81; DB 20; Length 15; 100.0%; Pred. No. 3.2e-06; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                              Immunodominant epitope from p53 N-terminal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              transformation-associated protein.
                                                                                                                                                                                                                                                                                                     AARS4909 standard; peptide; 15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Legros Y, Lubin R, Soussi T;
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nes 14; Conservative
                                                                                                                                              Best Local Similarity 100.
Matches 15; Conservative
                                                                                                                                                                                             2 PLSQETFSDLWKLLP 16
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                                                                                                      15 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens.
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                                                                     neoplasia.
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                                                                                                      Sequence
                                                                                                                                   Query Match
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                                                                                                                                                                                                                                                                          RESULT 9
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AAR54910

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Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptides derived from the N-terminal (amino acids 1-112) or the C-terminal (amino acids 350-393) of protein p53 which specifically react with anti-1953 antibodies in patients with cancer or precancerous conditions are claimed. The peptides (AARS4907-RS4921) are useful for detecting and monitoring cancerous and precancerous
                                                                                                 cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen; transformation-associated protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New immuno:dominant epitope(s) of protein p53 - for detecting and monitoring antibodies indicative of cancer and precancerous
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                                                                   Immunodominant epitope from p53 N-terminal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR89914 standard; peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 5; Page 42; 62pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                               Lubin R, Soussi T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95WO-GB01719.
                                                                                                                                                                                                                                                                                                  93WO-FR01082.
                                                                                                                                                                                                                                                                                                                                        92FR-0013110.
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                                   (first entry)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            15 AA;
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                                   29-NOV-1994
                                                                                                                                                                                        Homo sapiens,
                                                                                                                                                                                                                                                                                                    02-NOV-1993;
                                                                                                                                                                                                                                                                                                                                        02-NOV-1992;
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                                                                                                                                                                                                                                                              11-MAY-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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AAR54910;
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p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein identified by modifying the p53 consensus binding sequence by substitution of an Alanine at each pos. and identifying which and acid changes altered binding to expressed MDM2. The peptides and methods of identifying aniliar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New peptides of the invention which interfere with the binding of the
                                                                                                                                                                                                                                                                                                                        New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and other malignancies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Disclosure; Page 1; 46pp; English.
19-APR-1995; 95US-0424957.
                                   94US-0277660.
                                                                                                                                                                                Lane DP, Picksley SM;
                                                                                                     (UYDU-) UNIV DUNDEE.
                                                                                                                                                                                                                                                      WPI; 1996-105905/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15 AA;
                               20-JUL-1994;
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/ Match 73.3%; Score 77; DB 17; Length 15; Local Similarity 100.0%; Pred. No. 1.4e-05; les 14; Conservative 0; Mismatches 0; Indels 5 OETFSDLWKLLPEN 18 1 OETFSDLWKLLPEN 14 Query Match Matches δ

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0; Gaps

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Fork head associated; FHA; domain; transcriptional control; DNA replication; DNA repair; cell cycle control.
           AAB29157 standard; Peptide; 15 AA.
                                                                                                                  17-MAR-2000; 2000WO-GB01024.
                                    (first entry)
                                                                                          WO200057184-A2.
                                                                             Unidentified
                                   02-FEB-2001
                                                                                                      28-SEP-2000.
                                                Peptide #1.
                       AAB29157;
RESULT 12
      AAB29157
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(KUDO-) KUDOS PHARM LTD. WPI; 2000-664872/64. Jackson SP,

99GB-0015075.

99GB-0006432.

19-MAR-1999;

28-JUN-1999;

Assays and screening methods based on direct interaction between FHA domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide - buipuiq

Disclosure; Fig 2; 92pp; English.

The present invention describes isolated prepared P53 epitopes (I). Also described are: (1) a cytotoxic T lymphocyte (CTL) that is cultured in vivo and binds to a complex of (I); (2) a peptide (II) comprising (I) and a second epitope and has less than 50 contiguous amino acids, (3) a vaccine composition comprising (II), a unit dose of a peptide with less sequence of p53, and a pharmaceutical excipient; (4) an isolated nucleic acid encoding (I); and (5) an isolated nucleic acid encoding (I); and (5) an isolated nucleic acid encoding (II). (I) as cytostatic activity and can be used in vaccines. The vaccine composition is useful for treating or preventing cancer. (I) and (II) are useful as diagnostic agents and for evaluating immune responses. Unlike conventional epitopes, immunosuppressive epitopes that may be The present invention relates to assays and screening methods based on a direct interaction between fork head associated (FHA) domains and phosphorylated polypeptides, for characterizing the binding of these molecules. FHA peptides may be useful for treating medical conditions associated with defects in transcriptional control, DNA replication, DNA repair, cell cycle control or other cellular process. The method may provide valuable insights into checkpoint signalling, has important implications for the functions of other FHA domain-containing proteins and provides basis for new lines of therapy. The present sequence is a peptide used in the present invention. Gaps Epitope-based vaccines comprising P53 epitope having a specified sequences, useful for treating and preventing cancer, the epitopic peptides is useful as diagnostic agents and for evaluating immune Human; p53; tumour suppressor; human leukocyte antigen; HLA; CTL; cytotoxic T lymphocyte; T cell; immune response; tumour; cancer; vaccine; epitope; cytostatic. Celis E; 0; Length 15; Indels Southwood S, Chesnut R, p53 DR 3a motif binding peptide exemplary sequence #3. 73.3%; Score 77; DB 21; Le 100.0%; Pred. No. 1.4e-05; tive 0; Mismatches 0; Disclosure; Page 113; 138pp; English. AAG89500 standard; Peptide; 15 AA. Sette A, Sidney J, 11-DEC-2000; 2000WO-US33629. 99US-0458297 Query Match
Best Local Similarity 100.0
Matches 14; Conservative 11-SEP-2001 (first entry) 1 PPLSOETFSDLWKL 14 2 PPLSOETFSDLWKL 15 (EPIM-) EPIMMUNE INC. WPI; 2001-381493/40. 15 AA; WO200141788-A1. 10-DEC-1999; Homo sapiens. 14-JUN-2001 Synthetic. Seguence AAG89500; Fikes J, Keogh E; response RESULT 13 AAG89500 8*66666666666 à В

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The present invention describes isolated prepared P53 epitopes (I). Also described are: (I) a cytotoxic T lymphocyte (CTL) that is cultured in vivo and binds to a complex of (I); (2) a peptide (II) comprising (I) and a second epitope and has less than 50 contiguous amino acids; (3) a vaccine composition comprising (II), a unit dose of a peptide with less than 50 contiguous amino acids with 100% identity to the native peptide sequence of p53, and a pharmaceutical excipient; (4) an isolated nucleic acid encoding (I); and (5) an isolated nucleic acid encoding (II). (I) acid encoding (I); and (5) an isolated nucleic acid encoding (II). (I) are useful as diagnostic agents and for evaluating immune responses. Composition is useful for treating or preventing immune responses. Unlike conventional epitopes, immunosuppressive epitopes that may be present in whole antigens can be avoided with the use of the vaccine composition of (I). The ability to combine selected epitopes and
                Composition of (1). The ability to combine selected epitopes and further, to modify the composition of the epitopes enhances the immunospanicity. The possible pathological side effects caused by infectious agents or whole pathological subjections agents or whole primarises, which might have their own intrinsic biological activity, are eliminated. AAG89363 to AAG89747 represent amino acid sequences used in the exemplification of the
                                                                                                                                                                                                                                                              Gaps
present in whole antigens can be avoided with the use of the vaccine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Epitope-based vaccines comprising P53 epitope having a specified sequences, useful for treating and preventing cancer, the epitopic peptides is useful as diagnostic agents and for evaluating immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human, p53, tumour suppressor, human leukocyte antigen, HLA, CTL, cytotoxic T lymphocyte, T cell, immune response, tumour, cancer, vaccine, epitope, cytostatic.
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                                                                                                                                                                                                                      73.3%; Score 77; DB 22; Length 15; 100.0%; Pred. No. 1.4e-05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                              AAG89730 standard; Peptide; 15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    p53 DR3 binding peptide #1.
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                                                                                                                                                                                                                                                                                                    1 PPLSQETFSDLWKL 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2001-381493/40.
                                                                                                                                                                                                                                           Best Local Similarity
                                                                                                                                          present invention.
                                                                                                                                                                                15 AA;
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Synthetic.
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Keogh E;
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          immunogenicity. The possible pathological side effects caused by infectious agents or whole protein antigens, which might have their own intrinsic biological activity, are eliminated. AAG89363 to AAG89747 represent amino acid sequences used in the exemplification of the
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                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                 Immunisation, immunogenicity enhancer, rabies virus strain ERA; immunodominant T-helper cell determinant; nucleoprotein; antibody; post-translational modification; p53; cancer; diagnosis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Methods of detecting post-translational modification level of p53. useful for diagnosis of cancer
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further, to modify the composition of the epitopes enhances the
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                                                                                                                               73.3%; Score 77; DB 22; Length 15; 100.0%; Pred. No. 1.4e-05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Disclosure; Page 45; 59pp; English.
                                                                                                                                                                                                                                                                                               AAY45227 standard; peptide; 17 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Thurin M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US04653.
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Search completed: February 12, 2003, 11:13:44 Job time: 47.5 secs

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Copyright (c) 1993 - 2003 Compugen Ltd.
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Run on:

February 12, 2003, 11:12:02; Search time 37.2083 Seconds (without alignments) 105.216 Million cell updates/sec

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19

Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

671580 segs, 206047115 residues Searched:

Total number of hits satisfying chosen parameters:

9297

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SPIREMBL 21:* Database :

sp archea:*
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SUMMARIES

					COLUMNITES	
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Result		Query	Query			
No.	Score	Match	Length	DB	ID	Description
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	PRT;	Created)	Last seque										P.A., Green	Sibson D.R., Vergote I., Walker	EMBL/GenBank/DDBJ databases.		;	•			·	•	•					D7FA272EBFAB9798 CRC64;	Score 77;	Pred. No. 1.	
	PRELIMINARY;	(TrEMBLrel, 15,	(TrEMBLrel. 15,	(TrEMBLrel. 17,	Tumor suppressor p53 (Fragment).		(Human).		theria; Primates;			N.A.	Thompson-Hehir J., Davies M.P.A., Green J.A., Halliwell N.	Joyce K.A., Salisbury J., Si		6362.1;	28; AAF36354.1;							35; AAF36361.1;); P53; 1.	25 25	25 AA; 2890 MW;		100.0%;	
JLT 1 168	Q9NP68 Q9NP68;	01-OCT-2000		01-JUN-2001	Tumor suppre	TP53.	Homo sapiens (Human).	Eukaryota; Me	Mammalia; Eutheria;	NCBI_TaxID=9606;	[]	SEQUENCE FROM N.A.	Thompson-Heh	Joyce K.A., 5	Submitted (NOV-1999)	EMBL; AF209136;					EMBL; AF209132;			EMBL; AF209135;	InterPro; IPR002117;	Pfam; PF00870; P53;	NON_TER	SEQUENCE 25	Query Match	Best Local Similarity Matches 14: Conserv	,
RESULT Q9NP68	A ID	DŢ	DŢ	БŢ	DE	S	SO	ပွ	ပ	ŏ	Z.	RP	RA	ΚA	RL	DR	DR	DR	DR	DR	DR	DR	DR	DR	DR	N.	БŢ	SO	8	M K	1

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           Q85618
Q85618;
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085618
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                                                                                                                                                                                       SEQUENCE FROM N.A.
MEDLINE=99067015; PubMed=9851690;
Grechez-Cassiau A., Bernard M., Ladjali K., Rodriguez I.R., Voisin P.;
"Structural analysis of the chicken hydroxyindole-O-methyltransferase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   dallus gallus (Chicken).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Archosauria; Aves; Neognathae; Galliformes; Phasianidae; Phasianinae;
                                                                                                           Gallus gallus (Chicken).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Archosauria, Aves, Neognathae, Galliformes, Phasianidae, Phasianinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIJNE=84144799; PubMed=6322173; Westaway D., Payne G., Varmus H.E.; "Proviral deletions and oncogene base-substitutions in insertionally mutagenized c-myc alleles may contribute to the progression of avian bursal tumors.";
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                                                                                                                                                                                                                                                                                                                                      / Match 42.9%; Score 45; DB 13; Length 22; Local Similarity 58.3%; Pred. No. 2.5; nes 7; Conservative 4; Mismatches 1; Indels
                                                                   01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Hydroxyindole-o-methyltransferase isoform A (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-UN-2002 (TrEMBLrel. 21, Last annotation update)
C-myc oncogene, aa 44-62 from c-myc/LL3 (ALV provirus)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE 24 AA; 2679 MW; 93F59F2D3B24E527 CRC64;
                                                                                                                                                                                                                                                                                                              SEQUENCE 22 AA; 2783 MW; 35726EABF1E450C8 CRC64;
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EMBL; K01440; AAA48964.1; -.
InterPro; IPR002418; TF_Myc.
PFam; PF01056; Myc_N term; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           24 AA.
                                                      01-NOV-1999 (TrEMBLrel. 12, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                              PRT;
                                                                                                                                                                                                                                                          Eur. J. Biochem. 258:44-52(1998).
                                                                                                                                                                                                                                                                         EMBL; AF116455; AAD23444.1; -.
Methyltransferase; Transferase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 PPAPSE---DIWKKFELLP 17
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                              PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                               4 SQETFSDLWKLL 15
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                                                                                                                                                               NCBI_TaxID=9031;
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                                                                                                                                                     Gallus.
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                              CO9M6D
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    RESULT 2
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               Q9W6D7
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-!- FUNCTION: THIS PROTEIN BINDS CALCIUM (BY SIMILARITY).
-!- SINGELLULAR LOCATION: BNDOPLASMIC RETCULUM LUMEN.
-!- SINGELLULAR LOCATION: BNDOPLASMIC RETCULUM LUMEN.
-!- SINGELLULAR LOCATION: TROPHOZOITES.
-!- SINGELLULAR LOCATION: TROPHOZOITES.
-!- SINGELLULAR LOCATION: IN TROPHOZOITES.
-!- SINGELLULAR LOCATION: I. PARTIAL.
--- PROSITE: PSO0804; CALRETICULIN 2: PARTIAL.
--- PROSITE: PSO0804; CALRETICULIN 2: PARTIAL.
--- PROSITE: PSO0804; CALRETICULIN 2: PARTIAL.
--- PROSITE: PSO0805; CALRETICULIN 2: PARTIAL.
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                                                                                                                                                                                                                                                                                                                                                             MEDLINE-84144799; PubMed-6322173; Westeaway D., Payne G., Varmus H.E.; Westeaway D., Payne G., Varmus H.E.; "Proviral deletions and oncogene base-substitutions in insertionally mutagenized c_myc alleles may contribute to the progression of avian
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ö
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                                          01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 21, Last annotation update)
RAV-2 proviral DNA, c-myc (Aa 44-67), allele LL3 (Fragment).
AAV-3 provirus.
Viruses, Retroid viruses; Retroviridae; Alpharetrovirus.
NCBI_TaxID=11864;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31.4%; Score 33; DB 15; Length 24; 47.4%; Pred. No. 2.40+02; trive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  30.5%; Score 32; DB 5; Length 19; 45.5%; Pred. No. 2.8e+02; trive 3; Mismatches 3; Indels
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01-0CT-2001 (TrEMBLrel. 18, Last sequence update)
01-0CT-2001 (TrEMBLrel. 18, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Acad. Sci. U.S.A. 81:843-847(1984).
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Entamoeba histolytica.
Bukaryota, Entamoebidae, Entamoeba.
NCBI_TaxID=5759;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
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EMBL; K01439; AAA51600.1; -.
InterPro; IPR002418; TF Myc.
Pfam; PF01056; Myc. N. term; 1.
NON TER 1 24 24
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Matches 9; Conservative
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PRELIMINARY;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       bursal tumors."
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1 PPLSQETFSD 10
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NON TER
SEQUENCE
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MEDLINE=20248365; PubMed=10789503;
Chida T., Okamura N., Ohtani K., Yoshida Y., Arakawa E., Watanabe H.;
"The complete DNA sequence of the O antigen gene region of Plesiomonas shigelloides serotype 017 which is identical to Shigella sonnei form
                                                                                                                                                                                                                                                                                                                               Houng H.H., Venkatesan M.M., "Genetic analysis of Shigella sonnei form I antigen: identification of a novel IS630 as an essential element for the form I antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Larkin D., Serov O., Zhdanova N.; Mapping of five genes from human chromosome 17 to chromosome hn of the acompon shrew (Sorex araneus)...; Acta Theriol. (Warsz) 45:143-146(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sorex araneus (Eurasian common shrew) (European shrew).
Wakaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Insectivora; Soricidae; Soricinae; Sorex.
                                                                                                                 Shigella sonnei.
Bacteria, Proteobacteria, gamma subdivision, Enterobacteriaceae,
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Pred. No. 3.7e+02;
1: Mismatches 7; Indels
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                                                                                                                                                                                                                                                                                                                                                                             Microb. Pathog. 25:165-173(1998).
EMBL; A8028134; BAA85070.1; -.
NON TER 25 25
SRQÜENCE 25 AA; 2726 MW; A6CC5D24211CDAAD CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  16 AA; 1951 MW; 775186E3FE5F52E2 CRC64;
                                                                               Last sequence update)
Last annotation update)
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01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
                                              25 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Thyroid hormone receptor alpha (Fragment). THRA1.
                                                                                                                                                                                                                                                                  Microbiol. Immunol. 44:161-172(2000).
                                                                     Created)
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                                                                                                                                                                                                                                                                                                                     MEDLINE=99036814; PubMed=9817819;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACCA TREFIOL. (WAISZ) %J:13-1.
EMBL; AF314827; AAK13419.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       42.98;
                                                                 01-MAY_2000 (TrEMBLrel. 13,
01-MAY_2000 (TrEMBLrel. 13,
01-DEC-2001 (TrEMBLrel. 19,
ORF10S (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                              30.5%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     6; Conservative
                                             PRELIMINARY;
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                                                                                                                                           Shigella.
NCBI_TaxID=624;
                                                                                                                                                                                                                                                                                                                                                                      expression.";
                                                                                                                                                                                                                                                                                                         STRAIN=HW383;
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SEQUENCE
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Bukaryota; Metazoa; Nematoda; Chromadorea; Rhabditida; Rhabditoidea;
Rhabditidae; Peloderinae; Caenorhabditis.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bos taurus (Bovine).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Cetartiodactyla, Ruminantia, Pecora, Bovoidea,
Bovidae, Bovinae, Bos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE=99175163; PubMed=10075683; Jiang H., Okamura C.S., Lucy M.C.; Jiang H., Okamura C.S., Lucy M.C.; Isolation and characterization of a novel promoter for the bovine growth hormone receptor gene."; J. Biol. Chem. 274:8933-7900(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ·.
                                                                                                                                                                                                                                                                                                                                                         Zeke T., Gergely P., Dombradi V.;
"The catalytic subunits of Ser/Thr protein phosphatases from
Caenorhabditis elegans: A biochemical and molecular biological
                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (JUL-1996) to the EMBL/GenBank/DDBJ databases.
-!- CATALYTIC ACTIVITY: A PHOSPHOPROTEIN + H(2)O = A PROTEIN PHOSPHATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         -!- SIMILARITY: BELONGS TO THE PPP FAMILY OF PHOSPHATASES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   29.5%; Score 31; DB 5; Length 24; 41.7%; Pred. No. 5.2e+02; tive 4; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2BF73DBBB5F20359 CRC64;
                                                                                                                                           01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Last sequence update)
Last annotation update)
                                                                                                                                                                                                            PPI-like Ser/Thr protein phosphatase (Fragment). Caenorhabditis elegans.
                                                                                                          24 AA
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HSSP; P08129; 1FJM.
InterPro; 1PR004844; S/T_phosphtse.
ProDom; PD000252; S/T_phosphtse; 1.
Hydrolase; Iron; Manganese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Growth hormone receptor (Fragment).
                                                                                                          PRT;
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01-NOV-1998 (TrEMBLrel. 08,
01-MAR-2001 (TrEMBLrel. 16,
01-MAR-2002 (TrEMBLrel. 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity 41.7
                                                                                                        PRELIMINARY;
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STRAIN=BREED HOLSTEIN;
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4 PPLFLEVFED 13
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PP1-like Ser/Thr protein phosphatase (Fragment).
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Q95V74;
                                                                                                                    survey."
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                                                                                                                                                                                                                                                                                                                  0; Gaps
                         Jiang H., Lucy M.C.; "Identification of 5' untranslated variants of bovine growth hormone
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Orlovsky I.V., Sverdlova P.S., Rubtsov P.M.; "Microsatellites within the human growth hormone receptor gene."; Submitted (JUL-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ295613; CAC06615.1; -
                                                                                                                                                                                                                                                                                          Match 28.6%; Score 30; DB 6; Length 14; Local Similarity 83.3%; Pred. No. 4.2e+02; les 5; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                / Match 28.6%; Score 30; DB 4; Length 23; Local Similarity 83.3%; Pred. No. 7.2e+02; Les 5; Conservative 1; Mismatches 0; Indels
                                                                                          Jiang H., Lucy M.C.;
"Bovine GH receptor 5' UTR variants.";
Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF036297; AAC33315.2;
                                                            Submitted (DEC-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NON TER 23 23
SEQUENCE 23 AA; 2426 MW; 9AD833A0F1DD699E CRC64;
                                                                                                                                                                                                                                                                    14 AA; 1544 MW; 5A45D458FF3A9E27 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-FBB-1997 (TrEMBLrel. 02, Created)
01-FBB-1997 (TrEMBLrel. 02, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                            01-WAR-2001 (TrEMBLrel. 16, Created)
01-WAR-2001 (TrEMBLrel. 16, Last sequence update)
01-WAR-2001 (TrEMBLrel. 16, Last annotation update)
Growth hormone receptor (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                       23 AA.
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                                                                                                                                      EMBL, AF085281; AAC22511] --
EMBL, AF036291, AAC33308.1; --
EMBL, AF036291, AAC33309.1; --
EMBL, AF036292; AAC33310.1; --
EMBL, AF036293; AAC33311.1; --
EMBL, AF036294; AAC33312.1; --
EMBL, AF036295, AAC33312.1; --
EMBL, AF036296; AAC33313.1; --
EMBL, AF036296; AAC33313.1; --
EMBL, AF326349; AAK97345.1; --
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               SEQUENCE FROM N.A.
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2 DLWQLL 7
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2 DLWQLL 7
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MEDLINE=99297341; PubMed=10368952;
Campbell N.J.H., Barker S.C.;
"The novel mitochondrial gene arrangement of the cattle tick,
The novel mitocoplus: fivefold tandem repetition of a coding region.";
Mol. Biol. Evol. 16:732-740(1999).
EMBL; AFILI0614; AAD28397.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
Caenorhabditis elegans.
Eukaryota, Metazoa, Nematoda, Chromadorea, Rhabditida, Rhabditoidea,
Rhabditidae, Peloderinae, Caenorhabditis.
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                                                                                                                                                                                                                                                    Submitted (JUL-1996) to the EMBL/GenBank/DDBJ databases.
--- CATALYTIC ACTIVITY: A PHOSPHOPROTEIN + H(2)O = A PROTEIN + PHOSPHATE.
--- SIMILARITY: BELONGS TO THE PPP FAMILY OF PHOSPHATASES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bukaryota, Metazoa, Arthropoda, Chelicerata, Arachnida, Acari,
Parasitiformes, Ixodida, Ixodidae, Boophilus.
                                                                                                                                                       Zeke T., Gergely P., Dombradi V.;
"The catalytic subunits of Ser/Thr protein phosphatases from
Caenorhabditis elegans: A biochemical and molecular biological
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27.6%; Score 29; DB 8; Length 17; 75.0%; Pred. No. 7.6e+02; tive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4; Indels
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NON TER 17 17
SEQUENCE 17 AA; 1988 MW; 319F2D4DA7DA11F3 CRC64;
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24 AA; 2781 MW; 353E423EA763AA75 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28.6%; Score 30; DB 5; L
58.3%; Pred. No. 7.5e+02;
tive 1; Mismatches 4;
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                                                                                                                                                                                                                                                                                                                                                      EMBL; Z77728; CAB01287.1; -.
HSSP; P08129; 1FJM.
INTERPRO; IPR004844; S/T phosphtse.
ProDom; PD000252; S/T phosphtse; Iydrolase; Iron; Manganese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Boophilus microplus (Cattle tick).
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Matches 7; Conservative
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                                                                             NCBI_TaxID=6239;
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Citrus sinensis (Sweet orange).
Bukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae;
NCBI_TaxID=2711;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                             Elongation factor 1 alpha (Fragment).

Whilus galloprovincialis (Mediterranean mussel).

Bukaryota, Metavoa; Mollusca, Bivalvia; Pteriomorphia; Mytiloida,
Mytiloidea; Mytilidae; Mytilus.
                                                                                                                                                         Bierne N., Daguin C., Borsa P., Jollivet D., Viard F., Bonhomme F.
David P.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Back J.-M., Kenerley C.M., The arg2 gene of Trichoderma virens: Cloning and development of a
                                                                                                                                                                                     "Introgression patterns in the Mytilus edulis / Mytilus
galloprovincialis mosaic hybrid zone along the Atlantic coasts of
                                                                                                                                                                                                                                                                                                                                                                     .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota, Fungi, Ascomycota, Pezizomycotina, Sordariomycetes,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 28.5; DB 3; Length 24;
Pred. No. 1.3e+03;
2; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                               27.6%; Score 29; DB 5; Length 24; 38.5%; Pred. No. 1.1e+03; Vative 3; Mismatches 5; Indels
                                                                                                                                                                                                                             France. , Submitted (SEP-2001) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       homologous transformation system.";
Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases
EMBL; AF001029; AAB58298.1; - SEQUENCE 24 AA; 2782 MW; 8FFD15C27EAD244C CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-JUL-1997 (TrEMBLrel. 04, Created)
01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Carbamoyl phosphate synthetase, small subunit (arg2).
Trichoderma virens.
                                                                                                                                                                                                                                                                                               SEQUENCE 24 AA; 2875 MW; 1D0DEAD3FBD13DCB CRC64;
01-DEC-2001 (TrEMBLrel. 19, Created)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-0CT-2000 (TrEMBLrel. 15, Created)
01-0CT-2000 (TrEMBLrel. 15, Last sequence update)
01-0CT-2000 (TrEMBLrel. 15, Last annotation update)
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NCBI_TaxID=29875;
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                                                                                                                                                                                                                                                   EMBL; AF424742; AAL24803.1; -.
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Matches 7; Conservative
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GenCore version 5.1.3
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Run on:

February 12, 2003, 11:11:31 ; Search time 10.2917 Seconds (without alignments) 76.572 Million cell updates/sec

Title: Perfect score: Sequence:

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

112892 segs, 41476328 residues Searched:

1520 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

SwissProt_40:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Description	P22702 neurospora	-		3 heliothis	9	Bpinac	5 acineto	P81626 locusta mig	1 sacchar	P80178 lymnaea sta	P80179 lymnaea sta	P80181 lymnaea sta	P80182 lymnaea sta	P24854 sus scrofa	P81490 phyllomedus	_	P04549 periplaneta		7	s	σ	Q9ur66 candida alb	S	P04548 periplaneta	P14596 tabanus atr		P82697 periplaneta	P82419 pachycondyl	P81217 equus cabal	٠.	0	_	9 lito
SUMMARIES	Y3KD NEUCR	LECB IRIHO	CRTC_SPIOL	HTF HELZE	CP35 PAPSP	TL17 SPIOL	1	AKHX_LOCMI	LPCP_YEAST	NP1_LYMST	NP2 LYMST	NP4_LYMST	NP5_LYMST	IBP4_PIG	DMS6_PHYBI	UR2_SCYCA	HTF2 PERAM	HTF2_CARMO	URE1 MORMO	ACT6_SOYBN	CPAX_BOVIN	NDK_CANAL	AKH_TABAT	HTF1_PERAM	HTF_TABAT	NP3_LYMST	PH4_PERAM	PCG6_PACGO				اريا	CR32_LITCE
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AC P36231:
DT 01-JUN-1994 (Rel. 29, Created)

DT 01-JUN-1994 (Rel. 29, Last sequence update)

DT 01-JUN-1994 (Rel. 29, Last annotation update)

DT 01-JUN-1994 (Rel. 29, Last annotation update)

BY 01-JUN-1994 (Rel. 29, Last annotation update)

RESULT 2

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litoria o sus scrot oryctola lactobaci treponem pachyconc pachyconc gryllus b thiobaci tenebrio nauphoete			; m	of the	. Fr.	th a collaboration EMBL outstation trictions on its is in no way und for commercial	1 1 1 1 1 4 4 4	1; Gaps		
P56240 P2644 P12664 P12664 P1983 P82424 P82424 P80408 P10919			Sordariomycetes	expression of ic carbamoyl	ODUCTION OF SE. 08521).	is produced through a collaid matics and the EMBL outs. There are no restrictions to as its content is in red. Usage by and for content the content is in red. Usage by and for content is the content is in the content in the content is in the content in the cont	4. 4	Length 24; Indels		
	TS.	update) nu update) 2 5'region.		. C.; s. Structure and ex f arginine-specific	S PR HETA	This SWISS-PROT entry is copyright. It is produced through a coll between the Swiss Institute of Bioinformatics and the EMBL out the European Bioinformatics Institute. There are no restriction use by non-profit institutions as long as its content is in modified and this statement is not removed. Usage by and for centities requires a license agreement (See http://www.isb-sib.ch/or send an email to license@isb-sib.ch).	8FF33F527EAD244C CRC64;	DB 1; e+02; 3;		
CR33_LITCE IRBP_PIG IRBP_RBIT UDP_LACCA FAGCA FAGCA PCW1_PACGO PCW2_PACGO PCW2_PACGO PCW2_PACGO PCW2_PACGO PCW2_PACGO PCW2_PACGO PCW2_PACGO PCW2_PACGO PCW2_PACGO PCW3_PACGO PCW3_PCW3_PACGO PCW3_PACGO PCW3_PCW3_PACGO PCW3_PACGO PCW3_PCW3_PACGO PCW3_PACGO PCW3_PCW3_PACGO PCW3_PACGO PCW3_PCW3_PACGO PCW3_PACGO PCW3_PCW3_PACGO PCW3_PACGO PCW3_PCW3_PACGO PCW3_PACGO PCW3_PCW3_PACGO PCW3_PACGO PCW3_PC	IGNMEN	Ω,	>- ·	5; cy C.; cus. Strr of argii	1990). FORY ROLI PHOSPHAT	OT entry is copyright. It is produ- Swiss Institute of Bioinformatics Bioinformatics Institute. There a profit institutions as long as i this statement is not removed. Us itres a license agreement (See http ail to license@isb-sib.ch).	F33F527E	Score 28.5; Pred. No. 2.8 ; Mismatches		
		Created) Last sequence Last annotation	ta; Pe Neur	141600 anofs) -2 loc bunit	D987 (: EGULA: AMOYL PA1 LI	copyright copyright ics Instit titutions ent is not nse agreem		0		
<u></u>		Creat Last Last	, co	I=2. Yargirgan	1 -1(RB)	tut tut iicg	MW.	₩ ₩		
00000000000000000000000000000000000000		19, 19, 40, CDa p	Ascomy ariacea	PubMed F M.S., rassa a	ase."; 5:10981 HAVE A IFIC CP	18-PROT entry is copy the Swiss Institute hean Bioinformatics I non-profit institut and this statement i requires a license in email to license@i	61 61 727	27.1%; larity 53.8%; Conservative	15	
0000000	į	(Rel. (Rel. (Rel. 2.8)	ssa 31; ord	N.A 33; ach; the	16t 26 1AY PEC		AAA336 B42224. protein	ity Ise	7 KE	
111111111111111111111111111111111111111		\sim \sim	crassa Fungi; ; Sordi	Se Se	mrth M: N M: N M: SE	S-PROT the Sw ean Bi non-pr and th requir n emai		lar Con	2-15 1-15 1-15 1-15 1-15 1-15 1-15 1-15	
00000000000000000000000000000000000000	T 1 NEUCR	P22702; P22702; 01-AUG-1991 01-AUG-1991 16-OCT-2001 Hypothetical	rospora aryota; dariales I_TaxID=	[1] SEQUENCE FROM N.A. MEDLINE=90293033; PubMed=2141606; Obbach M.J., Sachs M.S., Yanofsky C "The Neurospora crassa arg-2 locus. gene encoding the small subunit of a	phosphate synthetase "; J. Biol. Chem. 265:10981-10987(1990) -!- FUNCTION: MAY HAVE A REGULATORY ARGININE-SPECIFIC CARBAMOYL PHOS -!- SIMILARITY: TO YEAST CFA1 LEADER	This SWISS-PROT entry between the Swiss In the European Bioinform use by non-profit modified and this state entities requires a li or send an email to li	EMBL, J05512; PIR, B42224; B Hypothetical p SEQUENCE 24	Query Match Best Local Similarity Matches 7; Conser	4 SQETFSD-LWKLL : : 10 SQDYLSDHLWRAL	
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CP35_PAPSP
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          HTF_HELZE
RESULT 4
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                                                                     MEDLINE=93176159; PubMed=8439313;
Menegazzi P., Guzzo F., Baldan B., Mariani P., Treves S.;
"Purification of calreticulin-like protein(s) from spinach leaves.";
Blochem. Blophys. Res. Commun. 190:1130-1135(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Gaps
                                                                                                                                                                                                                                                0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                     Būkaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots,
Caryophyllidae, Caryophyllales, Chenopodiaceae, Spinacia.
Eukaryota; Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta; Magnoliophyta; Liliopsida; Asparagales; Iridaceae;
                                                               STRAIN-CV. GOLDEN HARVEST, and CV. PROF. BLAAUW; TISSUE-Bulb;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         24.8%; Score 26; DB 1; Length 20; 44.4%; Pred. No. 5.5e+02; tive 2; Mismatches 3; Indels
                                                                                                                                                                                                                        Score 27; DB 1; Length 20;
Pred. No. 3.9e+02;
1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -!- SUBCELLULAR LOCATION: Endoplasmic reticulum lumen.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -i- PTM: GLYCOSYLATED.
-i- SIMILARITY: BELONGS TO THE CALRETICULIN FAMILY.
PIR; PC1240; PC1241.
InterPro; IPR001580; Calreticulin.
PROSITE; PS00803; CALRETICULIN I; PARTIAL.
PROSITE; PS00804; CALRETICULIN 2; PARTIAL.
ENGORIES; PS00805; CALRETICULIN REPEAT; PARTIAL.
Endoplasmic reticulum; Calcium-binding; Glycoprotein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 AA; 2645 MW; 00FAB4C9DEEDCB0F CRC64;
                                                                                                                                                                                            20 20
20 AA; 2246 MW; DE08DA025FD17D56 CRC64;
                                                                                                                                                                                                                                                                                                                                                                 01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                             20 AA.
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                                                                                                                                                                                                                                                                                                                                           PRT;
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                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 44.4
Matches 4; Conservative
                                                                                                                                                                                                                                                                                                                                             STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                  Calreticulin (Fragment)
                                                                                                                                                                                                                                    Best Local Similarity
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                                  4CBI_TaxID=35876;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=3562;
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6 PLSEET 11
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SEQUENCE
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Heliothis zea (Corn earworm) (Bollworm).

Eukaryota, Metazoa, Arthropoda, Mandibulata, Pancrustacea, Hexapoda, Insecta, Pterygota, Neoptera, Endopterygota, Lepidoptera, Glossata, Ditrysia, Noctuoidea, Noctudae, Heliothinae, Helicoverpa.
                                                                                                                                                                    Papio sp. (Baboon).
Eukaryota, Metazoa; Chordata; Craniata, Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      01-WAY-1992 (Rel. 22, Created)
01-MAY-1992 (Rel. 22, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
Cytcohrome P450 3AS (EC 1.14.11) (CYPIIIAS) (P450 FA) (6-betatestosterone hydroxylase) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                     .
                                                                                                                                                                                                                                                                                                                                                                                           Score 24; DB 1; Length 10;
Pred. No. 5.3e+02;
                                                                                                                                                                                                                                                                                                                                           PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                                                                                         AMIDATION.
8E70367865A5B9D1 CRC64;
                      01-AUG-1990 (Rel. 15, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                     0; Mismatches
                                                           Hypertrehalosaemic hormone (HeZ-HRTH).
                                                                                                                                                                                                                                                                                                                                                                                            22.9%;
                                                                                                                                                                                                                                                                                                                                                                   10 AA; 1096 MW;
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                                                                                                                                                                                                                                                                                            PIR; A31571; A31571.
InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
Neuropeptide; Amidation.
MOD RES 10 10
                                                                                                                                                                                                                                                                                                                                                                                                                     5, Conservative
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STANDARD;
                                                                                                                                                         TISSUE=Corpora cardiaca;
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NCBI TaxID=61183;
                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 OLTFSSGW 8
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P80056;
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SEQUENCE
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Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots,
Caryophyllidae, Caryophyllales, Chenopodiaceae, Spinacia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     characterization.";
J. Biol. Chem. 273:6710-6716(1998).
J. Buol. Chem. 273:67100: Chloroplast; within the thylakoid lumen.
Chloroplast; Thylakoid.
NON TER
          PROSITE; PS00086; CYTOCHROME_P450; PARTIAL.
Oxidoreductase; Monooxygenase; Electron transport; Membrane; Heme;
Microsome; Endoplasmic reticulum.
NON TER 20 20
                                                                                                                                                                                                                                                                                                                                                  SEQUENCE, AND REVISIONS TO 13 AND 15.
Schubert M., Peterson U., Funk C., Schroeder W.P., Kieselbach T.;
"The chloroplast lumen from Arabidopsis thaliana.";
Submitted (JUL-2001) to the SWISS-PROT data bank.
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01-NOV-1995 (Rel. 32, Last sequence update)
16-OCT-2001 (Rel. 40, Last amontation update)
Benzaldehyde dehydrogenase [NAD+] I (EC 1.2.1.28) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Acinetobacter genomosp. 11.
Bacteria; Proteobacteria; gamma subdivision; Moraxellaceae;
Acinetobacter.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 24; DB 1; Length 23;
Pred. No. 1.3e+03;
1; Mismatches 1; Indels
                                                                             Score 24; DB 1; Length 20;
Pred. No. 1.1e+03;
                                                                                                   3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                              Andersson B., Schroeder W.P.;
                                                                                                                                                                                                                                           16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Thylakoid lumenal 17.4 kb protein (P17.4) (Fragment).
Spinacia oleracea (Spinach).
                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDINE-98175931; PubMed=9506969;
Kieselbach T., Hagman A., Andersson B., Schroeder W.F
"The thylakoid lumen of chloroplasts - isolation and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             23 AA; 2571 MW; 022C92717AE9AF2D CRC64;
                                                       SEQUENCE 20 AA; 2226 MW; 3FEF8B6E62BC0F36 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STRAIN=ATCC 11171 / NCIB 8250 / CIP 63.46 / B94;
MEDLINE=91113163; PubMed=1989592;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 23 AA
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                                                                                                      1; Mismatches
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InterPro; IPR001128; Cytochrome P450,
                                                                                                                                                                                                            PRT;
                                                                                                                                                                                                                                  Created)
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                                                                             22.9%;
                                                                                                    7; Conservative
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                                                                                                                             1 PPLSQETFSDLWKLL 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=106649;
                                                                                         Best Local Similarity
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                                                                                                                                                                                                                                                                                                                            NCBI TaxID=3562;
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P46365;
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SEQUENCE
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TL17_SPIOL
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"Comparison of benzyl alcohol dehydrogenases and benzaldehyde dehydrogenases from the benzyl alcohol and mandelate pathways in Acinetobacter calcoaceticus and from the TOL-plasmid-encoded toluene pathway in Pseudomonas putida. N-terminal amino acid sequences, amino acid compositions and immunological cross-reactions."; Biochem. J. 273:99-107(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Peptide hormone.
Locusta migratoria (Migratory locust).
Locusta migratoria (Migratory locust).
Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
Lusecta; Petrygota; Neoplera; Orthopteroidea; Orthopteroidea;
Acridomorpha; Acridoidea; Acrididae; Oedipodinae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Siegert K.J.;
Submitted (DEC-1998) to the SWISS-PROT data bank.
-!- FUNCTION: PROBABLY INVOLVED IN THE REGULATION OF LOCUST
-!- INTERMEDIARY WET-ABOLISM, BEHAVIOR AND/OR DEVELOPMENT.
-!- SIMILARITY: SOME SIMILARITY TO THE AKH / HRTH / RPCH FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ..
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-!- SIMILARITY: BELONGS TO THE ALDEHYDE DEHYDROGENASES FAMILY.
INTERPRO; IRR002086; Aldehyde dehydr.
PROSITE; PS00070; ALDEHYDE DEHYDR.CYS; PARTIAL.
PROSITE; PS00687; ALDEHYDE DEHYDR.GLU; PARTIAL.
Aromatic hydrocarbons catabolism; Oxidoreductase; NAD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      22.9%; Score 24; DB 1; Length 23; 23.1%; Pred. No. 1.3e+03; cive 3; Mismatches 7; Indels
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Pred. No. 7.6e+02;
0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                       23 AA; 2750 MW; 50496D4930DF9E40 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ol-AUG-1988 (Rel. 08, Created)
01-AUG-1988 (Rel. 08, Last sequence update)
01-NOV-1997 (Rel. 35, Last annotation update)
CPA1 leader peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       10 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Saccharomyces cerevisiae (Baker's yeast).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AMIDATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              62.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 62.33,
Best Local Similarity 62.33,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               InterPro; IPR002047; AKH. PROSITE; PS00256; AKH; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STANDARD;
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ID LPCP YEAST
AC P08521;
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Local Similarity
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P80179;
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P80181;
                                                                                                                                                           Query Match
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UNSURE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
                                                                                                                                                                                                                                                                                                           "Sequence and analysis of a 36.2 kb fragment from the right arm of yeast chromosome XV reveals 19 open reading frames including SNF2 (5'end), CPA1, SLY41, a putative transport ATPase, a putative ribosomal protein and an SNF2 homologue.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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Eukaryota, Metazoa, Mollusca, Gastropoda, Pulmonata, Basommatophora,
                                                                                                                                                    MEDLINE-85101411; Pubmed=3881260;
Werner M., Feller A., Pierard A.;
"Nucleotide sequence of yeast gene CP Al encoding the small subunit of arginine-pathway carbamoyl-phosphate synthetase. Homology of the deduced amino acid sequence to other glutamine amidotransferases.";

Eur. J. Biochem. 146:371-381(1985).
Eukaryota; Fungi; Ascomycota; Saccharomycotina; Saccharomycetes;
Saccharomycetales; Saccharomycetaceae; Saccharomyces.
NCBI_TaxID=4932;
                                                                                                                                                                                                                                                                                                                                                               Yeast 13:479-482(1997).
-!- SIMILARITY: TO A HYPOTHETICAL PROTEIN IN ARG2 5'REGION FROM NEUROSPORA CRASSA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1;
                                                 SEQUENCE FROM N.A.
MEDLINE=87215944; PubMed=3555844;
Werner M., Feller A., Messenguy F., Pierard A.;
The leader peptide of yeast gene CPAI is essential for the translational repression of its expression.";
Cell 49:805-813(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21.4%; Score 22.5; DB 1; Length 25; 50.0%; Pred. No. 2.4e+03; tive 2; Mismatches 2; Indels
                                                                                                                                                                                                                                                           STRAIN=S288c / FY1679;
BDDLINES-7298310; PubMed=9153758;
POITEY R., Cziepluch C., Tobiasch E., Pujol A., Kordes E., Jauniaux J.-C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              25 AA; 2979 MW; 01D494C8AFEAE4AA CRC64;
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01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
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EMBL; X01764; CAA25904.1; -.
EMBL; Z75210; CAA99620.1; -.
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nes 5; Conservative
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PIR; S25434; S25434.
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NCBL_TaxID=6523;
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TISSUE=Ganglion;
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SEQUENCE 25 A
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P80178;
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Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Basommatophora;
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Bukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Basommatophora;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Johnsen A.H., Rehfeld J.F.;
"LymnaDFamides, a new family of neuropeptides from the pond snail,
Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
MEDLINE=93238777; PubMed=8477756; Johnsen A.H., Rehfeld J.F., "LymuaDFamides, a new family of neuropeptides from the pond snail, LymnabFamides. Glue to cholecystokinin immunoreactivity in
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                                                                                                           Invertebraces?";

Eur. J. Biochem. 213:875-879(1993).
-!- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.

PIR; S32411; SA111.

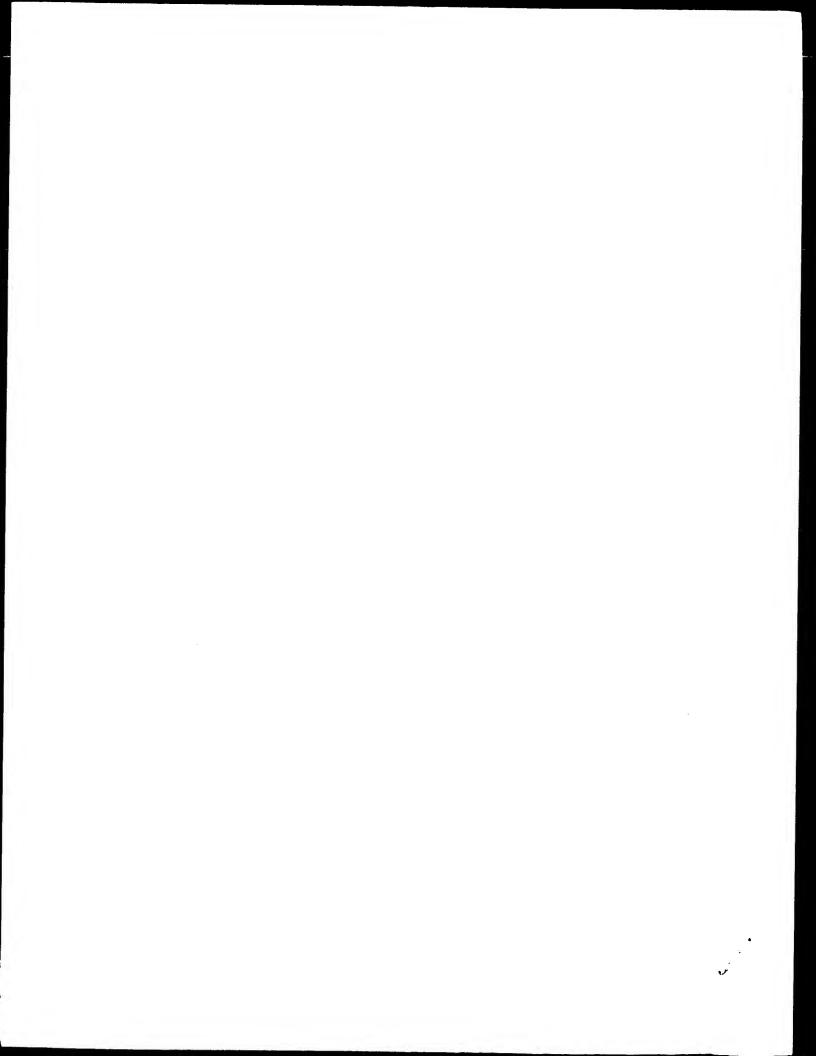
Neuropeptide, Amidation.

MOD_RES 13 AMIDATION.
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-!- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
PIR; S32472; S32472.
                                                                                                                                                                                                                                                                                                                                        21.0%; Score 22; DB 1; Length 13; 50.0%; Pred. No. 1.4e+03; tive 1; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                     9CA07BA3F5D5B455 CRC64;
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01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
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01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
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MEDLINE=93238777; PubMed=8477756;
                                                                                                                                                                                                                                                                                     SEQUENCE 13 AA; 1519 MW;
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MOD_RES 13 13
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Coleman M.E., Pan Y.-C.E., Etherton T.D.;

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Eukaryota, Metazoa, Mollusca, Gastropoda, Pulmonata, Basommatophora,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Ganglion;
MEDLINE=39238777; PubMed=8477756;
MDDINE=A.H., Rehfeld J.F.;
"LymnaDFanides, a new family of neuropeptides from the pond snail,
Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
MEDLINE=93238777; PubMed=8477756; Johnson A.H., Refield J.F., Johnson A.H., Refield J.F., Schner, Johnson A.H., Refield J.F., Inchange anides, a new family of neuropeptides from the pond snail, Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
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Mammalia, Butheria, Cetartiodactyla, Suina, Suidae, Sus.
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01-MAR-1992 (Rel. 21, Last sequence update)
15-DEC-1998 (Rel. 37, Last amnotation update)
Insulin-like growth factor binding protein 4 (IGFBP-4) (IBP-4)
(IGF-binding protein 4) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eur. J. Biochem, 213:875-879(1993).
-i- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
PIR; S32475; S32475.
Neuropeptide; Amidation.
MOD RES
MOD RES
                                                      invertebratës?";
Eur. J. Biochem. 213:875-879(1993).
-!- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 22; DB 1; Length 13;
Pred. No. 1.4e+03;
1; Mismatches 3; Indels
                                                                                                                                                            Length 13;
                                                                                                                                                                                       3; Indels
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                                                                                                                               9CA07BBB56D5B455 CRC64;
                                                                                                                                                            Score 22; DB 1; I
Pred. No. 1.4e+03;
                                                                                                                                                                                                                                                                                                                                                 01-JUL-1993 (Rel. 26, Created)
01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
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                                                                                                                                                                                       1; Mismatches
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50.0%;
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50.0%;
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                                                                                                    Neuropeptide, Amidation.
MOD RES 13 13
SEQUENCE 13 AA, 1503 M
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NCBI_TaxID=6523;
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hes 4; Conserv
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P80182;
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Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Phyllomedusinae; Phyllomedusa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "Structure, synthesis," and molecular cloning of dermaseptins B, a family of skin peptide antibiotics.";
J. Biol. (Dhen. 273:14680-14697(1998).
-:- FUNCTION: POSSESSES A POTENT ANTIMICROBIAL ACTIVITY AGAINST GRAM-POSITIVE AND GRAM-NEGATIVE BACTERIA. PROBABLY ACTS BY DISTURBING MEMBRANE FUNCTIONS WITH ITS AMPHIPATHIC STRUCTURE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE-98278974; PubMed=9614066; Charpentier S., Amiche M., Mester J., Vouille V., Le Caer J.-P., Nicolas P., Delfour A.;
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                                                                                                                                                                                                                                                    Length 16;
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-!- SIMILARITY: BELONGS TO THE DERMASEPTIN FAMILY.
Antibictic; Multigene family; Amphibian skin; Amidation.
MOD RES
SEQUENCE 24 AA; 2665 MW; E5987D7F50E08F4F CRC64;
                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                          SEQUENCE 16 AA; 1799 MW; 40988840096655E2 CRC64;
                                                                                                                                                                                                                                                 21.0%; Score 22; DB 1; I
66.7%; Pred. No. 1.8e+03;
tive 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                             15-DEC-1998 (Rel. 37, Created)
15-DEC-1998 (Rel. 37, Last sequence update)
16-OGT-2001 (Rel. 40, Last annotation update)
16-maseptin BVI (Dermaseptin B6)
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Phyllomedusa bicolor (Two-colored leaf frog).
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Best Local Similarity 50.v.
Local 4; Conservative
                                                                                                                                                                                                                                                                            4; Conservative
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                                                                                                                                                                                                                                                               Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=8393;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 LWKDILKN 9
                                                                                                                                                                                                                                                                                                                             7 PPPSEE 12
                                                                                                                                                                                                                                                                                                      1 PPLSQE 6
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                                                                                                                                                                                                                                                                                                                                                                                          DMS6 PHYBI
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                                                                                                                                                                                                                                                      Query Match
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DMS6_PHYBI
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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February 12, 2003, 11:12:22 ; Search time 20.5833 Seconds (without alignments) 88.739 Million cell updates/sec Run on:

US-09-403-440A-2

105 1 PPLSQETFSDLWKLLPENG 19 Perfect score: Seguence:

Gapop 10.0 , Gapext 0.5 **BLOSUM62** Scoring table:

Total number of hits satisfying chosen parameters: 283224 seqs, 96134422 residues Searched:

4984

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

PIR 73:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

genome polyprotein c-myc protein - ch protein phosphatas hypothetical prote gamma-crystallin I cytochrome P450 3A antifungal 2S stor Ig heavy chain DJ hypothetical prote Tha p 1 - Thaumeto hypertrehalosemic/ chaperonin 10 homo adipokinetic hormo adipokinetic hormo calcium-binding pr alpha-1-antitrypsi translation elonga pollen allergen Po ribulose-bisphosph testosterone 6beta base nonspecific a insulin-like growt benzaldehyde dehyd phosphoprotein pho lectin chain B - I H+-exporting ATPas (cytosine-5-)water-soluble 35K Ig heavy chain DJ Description SUMMARIES PC1240 A61506 PH1334 T42442 B42224 A35105 PA0110 A53592 A34921 PH1326 A31571 S21176 PC2347 D42842 S35641 S11545 A24244 A60372 S41747 S13298 T42259 В Query Match Length 30 Score Result

RESULT 3

T42442

protein kinase (EC	cytochrome-c oxida	phosphoprotein pho	gene CPA1 leader p	endo-1.4-beta-xvla	hypothetical prote	JymnaDFamide 1 - g	lymnaDFamide 2 - g	lymnaDFamide 4 - g	1 vmnaDFamide 5 - 3	1-2.4-diaminobutvr	ingulin-like growt	hydroxymroling-rio	njasma proteinase	intracrystalline o	nucleoside-diphosp
S20410	S65379	T42257	\$25434	870615	S16324	S32471	S32472	S32474	S32475	B44854	JH0517	S59481	A60812	A56894	S68023
7	7	7	N	~	~	N	~	7	~	7	7	~	N	7	7
15	23	24	25	S	60	13	13	13	13	14	16	17	20	20	21
21.9	21.9	21.9	21.4	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0	21.0
23	23	23	22.5	22	22	22	22	22	22	22	22	22	22	22	22
30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45.

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Genome polyprotein (clone L3/S2) - Skalica virus (fragment)
C;Species: Skalica virus
C;Species: Skalica virus
C;Species: Je-Mar-1997 #sequence_revision 19-Dec-1997 #text_change 17-Nov-2000
C;Accession: B61409
R;Guirakhoo, F.; Heinz, F.X.; Mandl, C.W.; Holzmann, H.; Kunz, C.; Gresikova, M.
A;Gen. Virol. '2, 333-338, 1991
A;Title: The relationship between the flaviviruses Skalica and Langat as revealed by mon A;Reference number: A61409; MUID:91132129; PMID:1847173
A;Accession: B61409
A;Accession: B61409
A;Molecule type: genomic RNA
A;Residues: 1-19 <GUI>C;Superfamily: yellow fever virus genome polyprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      R;Westaway, D.; Payne, G.; Varmus, H.E.
Proc. Natl. Acad. Sci. U.S.A. 81, 843-847, 1984
A;Title: Proviral deletions and oncogene base-substitutions in insertionally mutagenized
A;Reference number: I50387; MUID:84144799; PMID:6322173
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7
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C;Species: Gallus gallus (chicken)
C;Date: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 21-Jul-2000
C;Accession: IS0387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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A;Molecule type: DNA
A;Residues: 1-24 <WES>
A;Cross-references: GB:K01440; NID:g212357; PIDN:AAA48964.1; PID:g212358
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 24;
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B61409
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datifungal 2S storage albumin large chain - radish (fragment)
C;Species: Raphanus sativus (radish)
C;Species: Raphanus sativus (radish)
C;Date: 27-Apr-1993 #sequence_revision 18-Nov-1994 #text_change 12-Apr-1995
C;Accession: D42842
R;Terras, F.R.; Schoofs, H.M.; De Bolle, M.F.; Van Leuven, F.; Rees, S.B.; Vanierleyden, A;Title: Chem. 257, 1531-15309, 1992
A;Title: Analysis of two novel classes of plant antifungal proteins from radish (Raphanu A;Reference number: A42842; MUID:92348373; PMID:1639777
                                                                                                                                                                                                                                                 submitted to the Protein Sequence Database, September 2001
A;Description: Isolation and characterization of a major allergen of the pine Frocession A;Reference number: A59396
A;Accession: A59396
A;Status: preliminary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C,Accession: A60728
R;Pineau, T.; Galtier, P.; Bonfils, C.; Derancourt, J.; Maurel, P.
B;Pineau, T.; Galtier, P.; Bonfils, C.; Derancourt, J.; Maurel, P.
Biochem. Pharmacol. 39, 901-909, 1990
A;Title: Purification of a sheep liver cytochrome P-450 from the P450IIIA gene subfamily A;Reference number: A60728; MUID:90179800; PMID:2310415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A,Gene: CYP3A
C;Superfamily: human cytochrome P450 CYP3A5; cytochrome P450 homology
C;Keywords: electron transfer; endoplasmic reticulum; heme; monooxygenase; oxidoreductas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Molecule type: protein
A;Residues: 1-20 <PIN>
C;Comment: This cytochrome P450 isozyme is a member of the P450IIIA family but is not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cytochrome P450 3A, troleandomycin-induced - sheep (fragment)
N;Contains: oxidoreductase (EC 1.-.-.)
C;Species: Ovis sp. (sheep)
C;Date: 14-May-1993 #sequence_revision 14-May-1993 #text_change 05-Mar-1999
                                                                                                                                           C;Spēcies: Thaumetopoea pityocampa
C;Date: 03-Jun-2002 #sequence_revision 03-Jun-2002 #text_change 03-Jun-2002
C;Accession: A59396
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               24.8%; Score 26; DB 2; Length 18; 38.5%; Pred. No. 1.2e+03; tive 3; Mismatches 5; Indels
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A;Note: sequence extracted from NCBI backbone (NCBIP:109925)
                                                                                                                Tha p 1 - Thaumetopoea pityocampa (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A,Experimental source: L5
A,Note: IgE-binding protein, major allergen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 24.85
Best Local Similarity 38.55
Matches 5; Conservative
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A;Molecule type: protein
A;Residues: 1-20 <TER>
                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: protein A; Residues: 1-18 < MON>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Accession: A60728
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                                                         RESULT 6
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                                                                                        C.Accession: T42442
R.Zeke, T.; Gergely, P.; Dombradi, V.
Submirted to the EMBL Data Library, July 1996
A.Description: The catalytic subunits of Ser/Thr protein phosphatases from Caenorhabditi
A.Reference number: Z22025
A.A.Cession: T42442
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C,Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 21-Jul-2000 C;Accession: B42224  
R;Orbach, M.J.; Sachs, M.S.; Yanofsky, C. J. Biol. Chem. 265, 10981-10987, 1990 A;Title: The Neurospora crassa arg-2 locus. Structure and expression of the gene encodin A;Reference number: A42224; MUID:90293033; PMID:2141606
                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: EMBL:Z77728; PIDN:CAB01287.1
C;Superfamily: phosphoprotein phosphatase; phosphoesterase core homology; phosphoprotein
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R,Chiou, S.H.; Chen, S.W.; Lo, T.B.
R;Chiou, S.H.; Chen, S.W.; Lo, T.B.
FEBS Lett. 209, 107-110, 1986
A;Title: The amino-terminal sequences of four major carp gamma-crystallin polypeptides
A;Reference number: A91369; WUID:87105907; PMID:3803569
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ô
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       hyporhetical protein (arg-2 5' region) - Neurospora crassa
C;Species: Neurospora crassa
C;Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 21-Jul-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gamma-crystallin IV - common carp (fragment)
C;Species: Cyprinus carpio (common carp)
C;Date: 02-Jun-1988 #sequence_revision 02-Jun-1988 #text_change 21-Nov-1997
                       C.Species: Caenorhabditis elegans
C.Species: 03-Dec-1999 #sequence_revision 03-Dec-1999 #text_change 02-Jun-2000
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A;Residues: 1-24 <ORB>
A;Cross-references: GB:J05512; NID:g168864; PIDN:AAA33610.1; PID:g168866
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 28.6%; Score 30; DB 2; Length 24; Best Local Similarity 58.3%; Pred. No. 4.2e+02; Matches 7; Conservative 1; Mismatches 4; Indels
protein phosphatase - Caenorhabditis elegans (fragment)
                                                                                                                                                                                                                                                                                       A; Status: preliminary; translated from GB/EMBL/DDBJ
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Best Local Similarity 53.0.
The Conservative
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Best Local Similarity 75.09
Matches, 6; Conservative
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A; Residues: 1-24 <ZEK>
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R;Wasserman, R.; Galili, N.; Ito, Y.; Reichard, B.A.; Shane, S.; Rovera, G.
J. Exp. Med. 176, 1577-1581, 1992
A;Title: Predominance of fetal type DJH joining in young children with B precursor lymph A;Reference number: PH1302; MUID:93094761; PMID:1460419
A;Accession: PH1334
A;Molecule type: DNA
A;Residues: 1-23 < WAS>
C;Superfamily: immunoglobulin V region; immunoglobulin homology
C;Keywords: heterotetramer; immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: mitochondrion Neurospora crassa

C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 07-Dec-1999

C;Accession: A35105

R;Saville, B.J.; Collins, R.A.

Cell 61, 685-696, 1990

A;Title: A site-specific self-cleavage reaction performed by a novel RNA in neurospora m

A;Reference number: A35105, MUID:90263093; PMID:2160856

A;Accession: A35105

A;Actus: preliminary; not compared with conceptual translation

A;Residues: 1-14 < SAV>

C;Genetics:
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                                                                                                                                                                                                                                                                                                                       ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   hypothetical protein - Neurospora crassa mitochondrion (fragment)
                                                                                                                                                                                                                                                            24.8%; Score 26; DB 2; Length 23; 33.3%; Pred. No. 1.6e+03; tive 2; Mismatches 6; Indels
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pollen allergen Poa-pI - Kentucky bluegrass (fragment)
C,Species: Poa pratensis (Kentucky bluegrass)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              23.8%; Score 25; DB 2; I 54.5%; Pred. No. 1.3e+03; iive 0; Mismatches 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C;Accession: PA0110
R;Kamo, M.; Kawakami, T.; Tsugita, A.
submitted to JIPID, March 1995
A;Reference number: PA0109
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Best Local Similarity 33,3%
....hes 4; Conservative
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Best Local Similarity 54.57
Local 6; Conservative
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Matches 5; Conservative
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A, Genetic code: SGC3
C, Keywords: mitochondrion
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Comp. Biochem. Physiol. B 100, 293-296, 1991
A.Title: Purification, characterization, and acute phase response of plasma alpha-1-anti
A.Reference number: A61506; MUID:92191572; PMID:1724745
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                                                                                                                                                                                                                                                                                                                               calcium-binding protein, 53K - spinach (fragment)
C;Species: Spinacia oleracea (spinach)
C;Apte: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 23-Aug-1997
C;Accession: PC1240; PC1241
R;Menegazzi, P.; Guzzo, F.; Baldan, B.; Mariani, P.; Treves, S.
Biochem: Biophys. Res. Commun. 190, 1130-1135, 1993
A;Title: Purification of calreticulin-like protein(s) from spinach leaves.
A;Reference number: PC1240; MUID:93176159; PMID:8439313
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       alpha-1-antitrypsin - golden hamster (fragment)
C;Species: Mesocricetus auratus (golden hamster)
C;Date: 15-Oct.1994 #sequence_revision 15-Oct.1994 #text_change 11-May-2000
C;Accession: A61506
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Ig heavy chain DJ region (clone C280-120) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 07-May-1999
C;Accession: PH1334
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                                                   Query Match 24.8%; Score 26; DB 2; Length 20; Best Local Similarity 45.5%; Pred. No. 1.4e+03; Matches 5; Conservative 3; Mismatches 3; Indels
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C; Superfamily: wheat alpha-amylase inhibitor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Noče: 55K form
C;Superfamily: calreticulin
C;Keywords: calcium binding; metal binding
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A, Residues: 1-20 <ME2>
A, Experimental source: leaf
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A,Residues: 1-20 <MEN>
A,Experimental source: leaf
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A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-20 <AME>
                                                                                                                                                         1 PPLSQETFSDL 11
                                                                                                                                                                                      8 PPLLQQCCNNL 18
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5 EERFEDGWE 13
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A;Accession: PC1241
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C;Date: 11-Feb-1993 #sequence_revision 11-Feb-1993 #text_change 09-Jun-2000
C;Accession: A60372; A33086
C;Extamoddoullah, A.K.H.
A.K.H. Allergy Appl. Immunol. 93, 371-377, 1990
A;Title: Two-dimensional gel electrophoretic analyses of Kentucky bluegrass and rye gras
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      H-exporting ATPage (EC 3.6.3.6) 24k chain - Thermus aquaticus (fragment)
C;Species: Thermus aquaticus
C;Species: Ol-Dec-1995 #sequence_revision Ol-Dec-1995 #text_change O3-Jun-2002
C;Accession: A53592
R;Yokoyama, K.; Akabane, Y.; Ishii, N.; Yoshida, M.
J. Biol. Chem. 269, 12248-12253, 1994
A;Title: Isolation of prokaryotic V-0V-1-ATPage from a thermophilic eubacterium Thermus A;Reference number: A53592; MUID:94216345; PMID:8163530
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C;Keywords: hydroxyproline; pollen
F;5,8/Modified site: hydroxyproline (Pro) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 23.8%; Score 25; DB 2; Length 20; Best Local Similarity 36.4%; Pred. No. 2e+03; Matches 4; Conservative 1; Mismatches 6; Indels
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23.8%; Score 25; DB 2; Length 20;
Best Local Similarity 46.2%; Pred. No. 2e+03;
Matches 6; Conservative 3; Mismatches 4; Indels
                                                                                                                                                        ence of Poa p I allergen.
A,Reference number: A60372; MUID:91317571; PMID:2101126
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7 LSQEVEAEIQALL 19
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A,Molecule type: protein
A,Residues: 1-20 <YOK>
C,Keywords: hydrolase
                                                                                                                                                                                                                                                  A; Molecule type: protein A; Residues: 1-20 < BKR>
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                                                                                                                                                                                                                      A; Accession: A60372
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GenCore version 5.1.3
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February 12, 2003, 10:43:13; Search time 30 Seconds (without alignments) 18.635 Million cell updates/sec Run on:

Title:

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19 Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

262574 seqs, 29422922 residues Searched:

127244 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 19

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:* Issued_Patents_AA:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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	Description	Segment 13 Samuel	2	4 4	-	20.	1	20.	S	21	21	23	Sequence 6. Appli	è	Sequence 6. Appli	2	9	Segmence 17. April	17	'n	45	46.	4 5	equence 46			equence 10	Sequence 15, Appl	
SUMMARIES	σī	US-09-081-975-13	US-08-277-660A-1	US-08-277-660A-4	US-08-424-957-1	US-08-424-957-20	US-09-035-686-1	US-09-035-686-20	US-08-277-660A-5	US-08-424-957-21	US-09-035-686-21	US-09-081-975-23	US-09-280-047-6	US-08-208-573B-6	PCT-US95-02856-6	US-09-081-975-22	US-08-277-660A-9	US-08-424-957-17	US-09-035-686-17	US-09-236-415-5	US-08-424-957-45	US-08-424-957-46	US-09-035-686-45	US-09-035-686-46	US-09-400-653A-7	US-09-248-061B-7	-08-277-	US-08-277-660A-15	
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do	Query Match I	85.7	73.3	73.3	73.3	73.3	73.3	73.3		70.5	70.5	70.5	٠	•	61.9	57.1	56.2	56.2	56.2	56.2	54.3	54.3	54.3	54.3	54.3	54.3	53.3	53.3	
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Sequence Seq	ETO, AND	,,0
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00000000000000000000000000000000000000	RESULT 1 US-09-081-975-13 Sequence 13, P Fatent NO. 64z GENERAL INFOR TITLE OF IN TORRESPENDE STATE: CORPUTER: COMPUTER: COMPU	Query Match Best Local S Matches 16

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RESULT 4
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Patent No. 5702908
GENERAL INFORMATION
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                                                                     Sequence 1, Application US/08277660A
Patent No. 5702908
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 73.3%; Score 77; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 3.8e-06; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400 CITY: San Francisco STATE: California STATE: California ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRIT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A-60244/WHD
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TORNEVAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-602
TELEPHONE: (415) 781-1989
TELEPHONE: (415) 781-1989
TELERX: 910 277299
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 aming acids
                        1 PPLSQETFSELWNLLPEN 18
1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RY: United States 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5 QETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 QETFSDLWKLLPEN 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: linear
US-08-277-660A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS
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                                                                                                              RESULT 2
US-08-277-660A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE:
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Gaps
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Patent No. 5770377

Patent No. 5770377

APPLICANT: Lanc, David W.

APPLICANT: Lanc, David W.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof

NUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS:

ADDRESSEE: Flehr, Hobbach, Teet, Albritton & Herbert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 73.3%; Score 77; DB 1; Length 15; Best Local Similarity 100.0%; Pred. No. 3.8e-06; Matches 14; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESTEE Flehr, Hobbach, Test, Albritton & Herbert STREET: Flour Embarcadero Center, Suite 3400 CITY: San Francisco STRTEE: California COUNTRY: United States ZIP: 94111-4818 COUNTRY: RABABLE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/424,957 FILING DATE: 19-APR-1995 CLASSIFICATION: PRIOR APPLICATION: PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: US 08/277,660 FILING DATE: 20-JUL-1994 ATTORNEY/AGENT INFORMATION:
                                                                                                  SCHWARE: Patentin Release #1.0, Version #1.30 SCHWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A FILING DATE: 20-UUL-1994
CLASSIFICATION: 514
ATTORNEY AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A-61228/WHD
                                                                                                                                                                                                                                                                                                                                  A-60244/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS_DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-1
TELECOMMUNICATION: INFORMATION:
                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: A-(TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             : (415) 781-1989
(415) 398-3249
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TELEX: 910 277299
INFORMATION FOR SEQ 1D NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
TELEEX: 910 27729
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2 PPLSQETFSDLWKL 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
US-08-277-660A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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GENERAL INFORMATION:
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Patent No. 15770377
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Pickeley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flahr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ;
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                                                                                                                            73.3%; Score 77; DB 1; Length 15; 100.0%; Pred. No. 3.8e-06; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              73.3%; Score 77; DB 1; Length 15; 100.0%; Pred. No. 3.8e-06; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Francisco
CITY: California
CONTRY: United States
ZIP: 94111-4187

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Dreger, Walter H.
REGISTATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-UL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 1, Application US/09035686
; Patent No. 6153391
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15 amino acids
    15 amino acids
                                                                                                                                                                        14; Conservative
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Matches 14; Conservative
                                                                                                                                                                                                                                           1 QETFSDLWKLLPEN 14
                                                                                                                                                                                                                   5 QETPSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2 PPLSQETFSDLWKL 15
                                                                TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: unknown
US-08-424-957-20
                                                                                                                                                  Best Local Similarity
                      TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                      RESULT 5
US-08-424-957-20
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TELEX: 9
                                                                         JUS-08-424-957-1
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LENGTH:
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Gaps
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Detein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION NUMBER: US/09/035,686
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: A-61228/WHD FELECOMMUNICATION INFORMATION: TELEPHONE: (415) 781-1989
TELEPHONE: (415) 388-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
ATTOCNEY/AGENT INPORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: California
COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 20, Application US/09035686; Patent No. 6153391; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (415) 398-3249
TELEEX: 910 277-299
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                             STREET: Four Embarcadero
CITY: San Francisco
STATE: California
COUNTRY: United States
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 QETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 QETFSDLWKLLPEN 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 7
US-09-035-686-20
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RESULT 6
US-08-27-660A-5
; Sequence 5, Application US/08277660A
; Patent No. 5702908
; GENERAL INFORMATION:
; APPLICANT: Lane, David P.
; TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
; TITLE OF INVENTION: Protein and Therapeutic Application Thereof
; TITLE OF INVENTION: Protein and Therapeutic Application Thereof
; TITLE OF INVENTION: Protein and Therapeutic Application Thereof
; CNRESPED ADDRESSES: 17
; ADDRESSES: Flour Embarcadero Center, Suite 3400
; TREET: Four Embarcadero Center, Suite 3400
; STREET: California
; COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 73.3%; Score 77; DB 4; Length 15; 100.0%; Pred. No. 3.8e-06; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
FILING DATE: 20-JUL-1994
CLASSIFICATION: 514
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                           CLASSIPICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
FILING DATE: 19-APR-1995
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: DESGRY, MAILET H.
REGISTRATION NUMBER: 24,190
REPERENCE/DOCKET NUMBER: 24,190
REPERENCE/OCKET NUMBER: A.61228/WHD
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 398-3249
TELEFAX: (415) 398-3249
TELEFAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 20: SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acids
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                      CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Dreger, Walter H. REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-6
TELECOMMUNICATION: TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELÉX: 910 277299
INFORMATION FOR SEQ ID NO: 5: SEQUENCE CHARACTERISTICS:
LENGTH: 15 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
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TELEPHONE: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches 14; Conservative
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Best Local Similarity
                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY:
US-09-035-686-20
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Sequence 21, Application US/08424957
Patent No. 5770377
GENERAL INFORMATION:
Patent No. 5770377
GENERAL INFORMATION:
PICKSIPS, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSER: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
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                                                                                     70.5%; Score 74; DB 1; Length 15; 93.3%; Pred. No. 1.2e-05; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          70.5%; Score 74; DB 1; Length 15; 93.3%; Pred. No. 1.2e-05; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957 FILING DATE: 19-APR.1995 CLASSIFICATION:
APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660 FILING DATE: 20-UUL-1994
ATTORNEY AGENT INTERNATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Dreger, Walter H.
REGISTATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-035-686-21
Sequence 21, Application US/09035686
Patent No. 6153391
GENERAL INFORMATION:
APPLICANT: Pickeley, Steven M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415) , ._.
TELEPHONE: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 15 amino acids
TYPE: amino acid
STRANDEDNESS:
                                                                                  Query Match
Best Local Similarity 93.33
Matches 14, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity 93.3
Matches 14; Conservative
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                                                                                                                                                                                 2 PLSQETFSDLWKLLP 16
                                                                                                                                                                                                                1 PLSQETFSGLWKLLP 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY: unknown
US-08-424-957-21
STRANDEDNESS:
                          ; TOPOLOGY:
US-08-277-660A-5
                                                                                                                                                                                                                                                                                                RESULT 9
US-08-424-957-21
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MOLECULE TYPE: peptide
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Batent No. 641979

GENERAL INFORMATION:
APPLICANT: Kaelin, William
APPLICANT: Acelin, William
APPLICANT: METHODS OF TREATMENT USING
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: MRTHODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSED: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0
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APPLICAMT: Lane, David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof VUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS:

ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400

CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ;
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                                                                                                                                                                                                                                                    ZIP: 94111-4187

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: Diskette
COMPUTER: US COMPATIBLE
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                                                                                                                                                                                                                                                             FLILMS ALDE:

CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REGISTRATION NUMBER: A-61228/WHD
TELEPHONE: (415) 781-1989
TELEPHONE: (415) 781-1989
TELEFRAX: 910 277299
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                  CITY: San Francisco
STATE: California
COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PLSQETFSGLWKLLP 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: unknown
US-09-035-686-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STRANDEDNESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Bo
STATE: M
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NS-09-280-047-6
Sequence 6, Application US/09280047
Sequence 6, Application US/09280047
Sequence 6, Application US/09280047
SEGNERAL INFORMATION:
APPLICANT: GOUBLI, SAID A.
TITLE OF INVENTION: QUANTITATION OF INDIVIDUAL PROTEIN
TITLE OF INVENTION: LINASE ACTIVITY
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: Ross & Stevens, S.C.
STREET: One South Pinckney Street, P.O. Box 2599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: Misconsin
COUNTRY: U.S.A.

ZITE: S1701-2599
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,047
FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION NUMBER: US 08/208,573
FILING DATE: 04-MAR-1994
ATTORNEY/AGENT INFORMATION:
NAMME: SARA, CHARLES S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 31506.006
TELECOMMUNICATION INFORMATION:
MATTORNEY/AGENT NUMBER: 31506.006
TELECOMMUNICATION INFORMATION:
               CORRELL AFFILTATION NUMBER: US/99/081,975
FILING DATE: 12-MAY-1998
CLASSIPFCATION: 435
FRICH APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Elsenstein, Romaid I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELECHOME: 617-345-6054
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TELEFAX: (608) 257-9175
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 16 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 15 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 70.5
Best Local Similarity 81.2
Matches 13; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2 PLSQETFSDLWKLLPE 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: anino acid
STRANDEDMESS: single
; TOPOLOGY: linear
US-09-081-975-23
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STATE: Wiscons
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TELEFAX:
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0
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                                       Score 65; DB 3; Length 15;
Pred. No. 0.00031;
1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Ross & Stevens, S.C.
STREET: One South Pinckney Street, P.O. Box 2599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 61.9%; Score 65; DB 4; Length 15; 84.6%; Pred. No. 0.00031; tive 1; Mismatches 1; Indels
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Sequence 6, Application PC/TUS9502856

GENERAL INFORMATION:
APPLICANT: Gouell, Said A

TITLE OF INVENTION: QUANTITATION OF INDIVIDUAL PROTEIN
TITLE OF INVENTION: RINASE ACTIVITY
NUMBER, OF SEQUENCES: 8

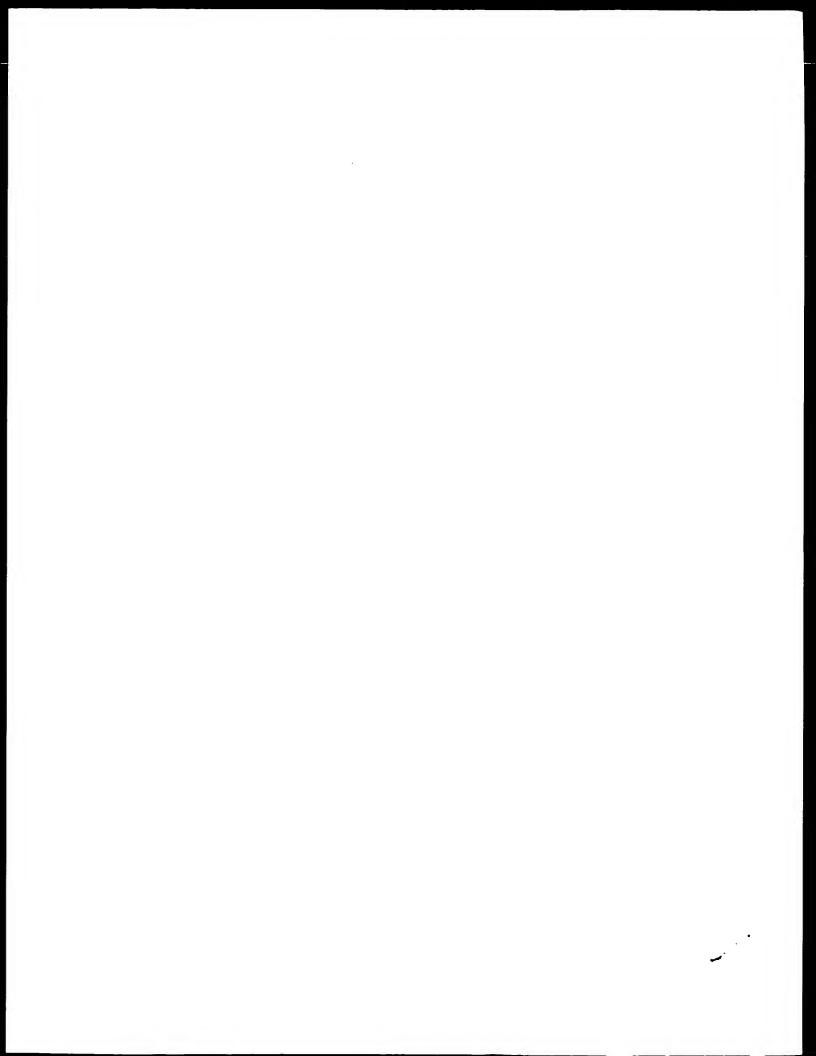
CORRESPENDENCE ADDRESS:
STREET: Bewitt Ross & Stevens, S.C.
STREET: Madison

CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DAIA:
APPLICATION NUMBER: US/08/208,573B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 10-Dec-1998
CLASSIFICATION: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: SARA, CHARLES S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUBER: 34506.006
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 257-5353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear ; MOLECULE TYPE: peptide ; SEQUENCE DESCRIPTION: SEQ ID NO: 6: US-08-208-573B-6
                                                                                                                                                                                                                                                                     Sequence 6, Application US/08208573B Patent No. 6348310 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CONTRY: U.S.A.
ZIP: 53701-2599
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (608) 257-9175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 15 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                          61.9%;
84.6%;
                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 10
                                   Query Match
Best Local Similarity 84.6'
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 84.6'
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Madison
                                                                                                                         1 PPLSQETFSDLWK 13
                                                                                                                                                  2 PPLSQEAFADLWK 14
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                                                                                                                                                                                                                             RESULT 13
US-08-208-573B-6
US-09-280-047-6
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APPLICANT: Kaelin, William
APPLICANT: Cost, Christine
TITLE OF INVENTION: METHODIES OF TREATMENT USING
TITLE OF INVENTION: METHODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
ADDRESSPRONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         61.9%; Score 65; DB 5; Length 15; 84.6%; Pred. No. 0.00031; tive 1; Mismatches 1; Indels
COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER FEADMABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IDM Compatible
COMPUTER: IDM Compatible
COMPUTER: BM Compatible
COMPETATING SYSTEM: Windows
SOFTWARE: FASTESQ for Windows
CURRENT APPLICATION DATA:
                                                                                                                                                                             APPLICATION NUMBER: PCT/US95/02856
                                                                                                                                                                                                                                                                                                                34506.006
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 22, Application US/09081975; Patent No. 6451979; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            47400
                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: SAZA, ChARLES S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 34500
TELEPHONE: 608-831-2100
TELEPHONE: 608-831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REPERENCE/DOCKET NUMBER: 474
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                9
                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 608-831-2100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 15 aming acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 84.6
Matches 11; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 PPLSQETFSDLWK 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 PPLSQEAFADLWK 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                amino acid
GY: linear
                                                                                                                                                                                                                         CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDALL
STREET: LV-
TW': Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                    FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: US
ZIP: 02110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PCT-US95-02856-6
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TELEK:

| TELEK:
| TELEK:
| INPORATION FOR SEQ ID NO: 22:
| SEQUENCE CHARACTERISTICS:
| LENGTH: 18 amino acids
| TYPE: amino a
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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Run on:

February 12, 2003, 10:44:28; Search time 56 Seconds (without alignments) 28.428 Million cell updates/sec

US-09-403-440A-2

105 1 PPLSQETFSDLWKLLPENG 19 Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

152190 Total number of hits satisfying chosen parameters:

425801 seqs, 83787487 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 19

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

Pending Patents AA New:*

1: /cgn2_6/ptodata/1/paa/USO6_NEW_COMB.pep:*

2: /cgn2_6/ptodata/1/paa/USO6_NEW_COMB.pep:*

3: /cgn2_6/ptodata/1/paa/USO8_NEW_COMB.pep:*

3: /cgn2_6/ptodata/1/paa/USO8_NEW_COMB.pep:*

5: /cgn2_6/ptodata/1/paa/USO8_NEW_COMB.pep:*

5: /cgn2_6/ptodata/1/paa/USO8_NEW_COMB.pep:*

7: /cgn2_6/ptodata/1/paa/USO8_NEW_COMB.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		de			SUMMARIES	
Result No.	Score	Query Match	Query Match Length	DB	ΙD	Description
-	92	87.6	19	-	US-60-438-805-9	Sequence 9. Appli
2	81	77.1	15	9	US-10-211-088-141	4
m	70	66.7	13	S	US-09-701-080C-22	
4	70	66.7	13	ß	US-09-701-080C-27	27,
S	59	56.2	11	ß	US-09-189-702A-286	286
9	41	39.0	14	Q	US-10-049-428-6	9
7	40	38.1	7	Ŋ	US-09-701-080C-12	12
œ	32	33.3	14	9	US-10-049-428-7	,
σ	31.5	30.0	14	9	US-10-268-332-53	Sequence 53. Appl
10	31.5	30.0	16	9	US-10-225-567A-1177	117
11	31	29.5	o,	Ŋ	US-09-189-702A-7	
12	31	29.5	12	Ŋ	US-09-281-717A-31	~
13	31	29.5	12	Ŋ	US-09-281-717A-33	i m
14	31	29.5	12	ß	US-09-281-717A-35	Seguence 35, Appl
15	31	29.2	12	S	US-09-830-693B-6	6.7
16	31	29.2	12	S	US-09-830-693B-8	
17	30	28.6	16	9	US-10-348-131-49	
18	29	27.6	15	9	US-10-169-026-3	'n
19	29	27.6	16	9	US-10-225-567A-942	
20	59	27.6	17	н	PCT-US02-28371-70	70.
21	29	~	17	9	US-10-236-878-70	70.
22	\sim	27.6	18	ß	US-09-552-802B-30	
23	28.5	27.1	18	ß	US-09-632-036E-11	-
24	28	26.7	7	9	US-10-174-717A-36	, 9
25	28	26.7	16	7	US-60-427-045-297	297
26	28	26.7	18	9	US-10-283-599-219	219,

TYPE: PRT

9, A 2575	Sequence 29042, A Sequence 25751, A	Sequence 29042, A Sequence 67, Appl	Sequence 18, Appl	463	144	Sequence 12880, A	Sequence 15661, A	Sequence 25791, A	Sequence 29085, A	463,	_	12880	15661	25791
000	9 6	US-09-641-528A-29042 US-10-029-206A-67	US-10-301-644-18 US-10-028-0758-67	US-09-641-528-463	US-09-641-528-1442	US-09-641-528-12880	US-09-641-528-15661	US-09-641-528-25791	US-09-641-528-29085	US-09-641-528A-463	US-09-641-528A-1442	US-09-641-528A-12880	US-09-641-528A-15661	US-09-641-528A-25791
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25.7		25.7	25.7	S	S	'n	25.7	25.7	25.7		25.7	25.7		25.7
28	27	27	27		27	27	27	27	27	27	27	27	27	27
23	300	31 32	33 4	35	36	37	38	39	40	41	42	43	44	4 5

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US-10-211-088-141

Sequence 14.1 Application US/10211088
GENERAL INFORMATION:
APPLICANT: Bright, Gary R.
APPLICANT: Premkumar, D. David
APPLICANT: Premkumar, D. David
TITLE OF INVENTION: NOVEL Fusion Proteins And Assays For Molecular Binding
FILE REFERENCE: 01-102-US
FILE REPERENCE: 01-102-US
CURRENT APPLICATION NUMBER: 05/10/211,088
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/349,395
PRIOR APPLICATION NUMBER: 60/341,589
PRIOR PILING DATE: 2001-12-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: Peptide US-60-438-805-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .;
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                     Sequence 9, Application US/60438805; GENERAL INFORMATION:
APPLICANT: KODADEK, THOMAS
ITILE OF INVENTION: MIXED BLEMENT CAPTURE AGENTS; TILE REFERENCE: UTSD:935USP1
CURRENT APPLICATION NUMBER: US/60/438,805; UNMBER OF SEQ ID NOS: 12; SOGTWARE: Patentin Ver. 2.1
SEQ ID NO 9; LENGTH: 19
                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 366
SOFTWARE: Patentin version 3.1
SEQ ID NO 141
LENGTH: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 PLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 PLSQETFSDLWKLLPEN 17
US-60-438-805-9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 56.2%; Score 59; DB 5; Length 11; Best Local Similarity 100.0%; Pred. No. 0.0046; Matches 11; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Xu, Weiming
APPLICANT: Liu, Lizhi
TITLE OF INVENTION: Unducible Screen for Drug Discovery
FILE REFERENCE: HO-P02380US0
CURRENT APPLICATION NUMBER: US/10/049,428
CURRENT APPLICATION NUMBER: 200-07-28
PRIOR APPLICATION NUMBER: GB 9918077
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                         APPLICANT: Sidney, John
APPLICANT: Kast, W. Martin
APPLICANT: Southwood, Scott
APPLICANT: Epimmune, Inc.
TITLE OF INVENTION: His Binding Peptides and Their Uses
FILE REFERENCE: 39963-20019.20
CURRENT APPLICATION NUMBER: US/09/189,702A
CURRENT FILING DATE: 1998-11-10
PRIOR PILING DATE: 1998-10-03-04
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  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: p53.14 peptide 34.0324
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SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                   ; Sequence 286, Application US/09189702A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 6, Application US/10049428; GENERAL INFORMATION: APPLICANT: Charles, Ian G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                               APPLICANT: Sette, Alessandro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 69.2
Matches 9; Conservative
  Matches 13; Conservative
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                                                           4 SQETFSDLWKLLP 16
                                                                                                               1 SQETFSDLWKLLP 13
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                                                                                                                                                                                           RESULT 5
US-09-189-702A-286
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US-10-049-428-6
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Sequence 22, Application US/09701080C

Sequence 22, Application US/09701080C

GENERAL INFORMATION:

APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY

TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION

FILE REPREMENCE: N7347C GGW

FILE REPREMENCE: N7347C GGW

FILE REPREMENCE: N7347C GGW

FILE REPREMENCE: N7347C GGW

FILE OF INVENTION: TRANSCRIPTIONAL REGULATION

FILE REPREMENCE: N7347C GGW

FILE OF INVENTION: TRANSCRIPTIONAL REGULATION

FRIOR PELLING DATE: 2001-02-27

FRIOR PELLING DATE: 1998-05-26

PRIOR PELLING DATE: 1998-01-05

NUMBER OF SEQ ID NOS: 36

NUMBER OF SEQ ID NOS: 36

SOFTWARE: PATENTI VET. 201-05
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN FITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN FITLE OF INVENTION: TRANSCRIPTIONAL RECULATION
FILE REPREMENT APPLICATION UNMERS: 05/09/701,080C
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: GB 9911303.8
PRIOR PLILING DATE: 1998-01-05
PRIOR APPLICATION NUMBER: GB 9900157.0
PRIOR APPLICATION NUMBER: GB 900157.0
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 27
FEMALE: 1190-01-05
FEMALE: PATENTIN VER. 2.1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence:derived from p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
                                                                                                                                                                                              0; Gaps
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                                                                                                                                        Query Match

77.1%; Score 81; DB 6; Length 15;
Best Local Similarity 100.0%; Pred. No. 2e-06;
Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 66.7%; Score 70; DB 5; Length 13; Best Local Similarity 100.0%; Pred. No. 9.6e-05; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 27, Application US/09701080C; GENERAL INFORMATION:
                                                              CTHER INFORMATION: Binding domain US-10-211-088-141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
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              ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                    1 PPLSQETFSDLWKLL 15
                                                                                                                                                                                                                                                       1 PPLSQETFSDLWKLL 15
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GENERAL INFORMATION:
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: LifeSpan Biosciences
APPLICANT: Burmer, Glenna C.
APPLICANT: Burmer, Glenna C.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
FILE REFERENCE: 1220-4-4
CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT APPLICATION NUMBER: 60/257,144
FRIOR APPLICATION NUMBER: 60/257,144
FRIOR PELING DATE: 2000-12-19
FRIOR FILING DATE: 2000-12-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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FILLE OF INVENTION: IMMUNE- AND COLON- RELATED TISSUES FILE REFERENCE: D0042A CIP
CURRENT APPLICATION NUMBER: US/10/268,332
CURRENT FILING DATE: 2002-10-10
PRIOR APPLICATION NUMBER: U.S. 60/235,713
PRIOR APPLICATION NUMBER: U.S. 60/261,783
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-01-16
PRIOR FILING DATE: 2001-07-13
PRIOR FILING DATE: 2001-07-13
PRIOR FILING DATE: 2001-08-17
PRIOR FILING DATE: 2001-08-17
PRIOR FILING DATE: 2001-08-17
PRIOR FILING DATE: 2001-09-26
NUMBER OF SEQ ID NOS: 66
SOFTWARE PARENT PARENT NOS: 66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
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GENERAL INFORMATION:
APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Bpimmune, Inc.
TITLE OF INFURION: HLA Binding Peptides and Their Uses
FILE REFERENCE: 39963-20019.20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
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Best Local Similarity 53.3.
8; Conservative
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5 SQENHS---RILPEN 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 7; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 53
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APPLICANT: Bristol-Myers Squibb Company
ITILE OF INVENTION: NOVEL HUMAN G-PROTEIN COUPLED RECEPTOR, HGPRBMY3, EXPRESSED HIGHI
                                              Sequence 12, Application US/09701080C
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: POLYPERIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 ITTLE OF INVENTION: TRANSCRIPTIONAL REGULATION
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT FILING DATE: 1998-05-26
PRIOR APPLICATION NUMBER: GB 9811303.8
PRIOR APPLICATION NUMBER: GB 9911303.8
PRIOR PILING DATE: 1998-05-26
PRIOR PILING DATE: 1998-01-05
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
NUMBER OF SEQ ID NOS: 36
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APPLICANT: Xu, Weiming
TITLE OF INVENTION Unducible Screen for Drug Discovery
TITLE OF INVENTION: Unducible Screen for Drug Discovery
FILE REPERENCE: HO-P02380US0
CURRENT APPLICATION NUMBER: US/10/049,428
CURRENT APPLICATION NUMBER: GB 9918077
PRIOR APPLICATION NUMBER: GB 9918077
PRIOR FILING DATE: 1999-07-30
PRIOR FILING DATE: 2000-06-30
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  33.3%; Score 35; DB 6; Length 14; 53.8%; Pred. No. 38;
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3; Mismatches
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Matches 7; Conservative
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Matches 7; Conservative
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                          US-09-701-080C-12
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LENGTH: 14
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APPLICANT: Kushner, Peter
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: Wagner, Richard
APPLICANT: Wagner, Richard
APPLICANT: Yamamoto, Keith
TITLE OF INVENTION: COACTIVACR BINDING
TITLE OF INVENTION: COACTIVACR BINDING
FILE REFERENCE: 9811-008-99
CURRENT APPLICATION NUMBER: US/09/281,717A
CURRENT FILING DATE: 1999-03-30
PRIOR PILING DATE: 1998-03-30
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.0
SEQ ID NO 31
LENGTH: 12
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29.5%; Score 31; DB 5; Length 9;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 6; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                           ) OTHER INFORMATION: p53.24V9 peptide 1323.02
US-09-189-702A-7
CURRENT APPLICATION NUMBER: US/09/189,702A
CURRENT FILING DATE: 1998-11-10
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1994-03-04
NUMBER OF SEQ ID NOS: 380
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 7
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OTHER INFORMATION: Leu --> Arg (L454R)
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OTHER INFORMATION: Leu --> Arg (L456R)
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; LOCATION: (8) .. (8)
; OTHER INFORMATION: Glu -->Lys (E457K)
US-09-281-117A-31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 31, Application US/09281717A; GENERAL INFORMATION:
, APPLICANT: Baxter, John
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Darimont, Beatrice
APPLICANT: Feng, Weijun
APPLICANT: Fletterick, Robert
                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 60.0%
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NAME/KEY: MUTAGEN
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US-09-281-717Ax33
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US-09-281-717A-31
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APPLICANT: Baxter, John
APPLICANT: Baxter, John
APPLICANT: Baxter, John
APPLICANT: Parimont, Beatrice
APPLICANT: Fletterick, Robert
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: West, Brian
APPLICANT: Wagner, Richard
APPLICANT: Yamamoto, Keith
TITLE OF INVENTION: COACTIVAOR BINDING
TITLE OF INVENTION: COACTIVAOR BINDING
FILE REFERENCE: 9811-008-99
CURRENT APPLICATION NUMBER: US 09/281,717A
CURRENT FILING DATE: 1999-03-30
PRIOR FILING DATE: 1998-03-30
PRIOR FILING DATE: 1998-03-30
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APPLICANT: Feng, Weijun
APPLICANT: Feng, Weijun
APPLICANT: Fleterick, Robert
APPLICANT: Weihner, Peter
APPLICANT: Wanner, Deter
APPLICANT: Wanner, Richard
APPLICANT: Wanner, Richard
APPLICANT: Yamamoto, Keith
TITLE OF INVENTION: COACTIVAOR BINDING
TITLE OF INVENTION: COACTIVAOR BINDING
TITLE OF INVENTION: US/09/281,717A
CURRENT APPLICATION NUMBER: US/09/281,717A
CURRENT APPLICATION NUMBER: US/09/281,717A
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29.5%; Score 31; DB 5; Length 12;

Best Local Similarity 60.0%; Pred. No. 1.4e+02;

Matches 6; Conservative 0; Mismatches 4; Indels
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PRIOR FILING DATE: 1998-03-30
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 35, Application US/09281717A; GENERAL INFORMATION:
APPLICANT: Baxter, John
Sequence 33, Application US/09281717A GENERAL INFORMATION:
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SOFTWARE: PatentIn version 3.0
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Best Local Similarity 60.v.
Local 6; Conservative
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US-09-281-717A-35
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US-09-830-693B-6
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LENGTH: 12
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LENGTH: 12
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## APPLICANT: Shiau, Andrew

APPLICANT: Shiau, Andrew

APPLICANT: Shiau, Andrew

APPLICANT: Rubiner, Peter

APPLICANT: Agard, David

APPLICANT: Agard, David

APPLICANT: Greene, Geoffrey

ITILB OF INVENTION: WETHODS AND COMPOUNDS FOR WODULATING NUCLEAR RECEPTOR

ITILB REPERRACE: 9311-013-99

CURRENT APPLICATION NUMBER: US/09/830,693B

FRIOR PILING DATE: 1999-03-30

PRIOR APPLICATION NUMBER: US 60/19,956

PRIOR APPLICATION NUMBER: US 60/113,146

PRIOR APPLICATION NUMBER: US 60/113,146

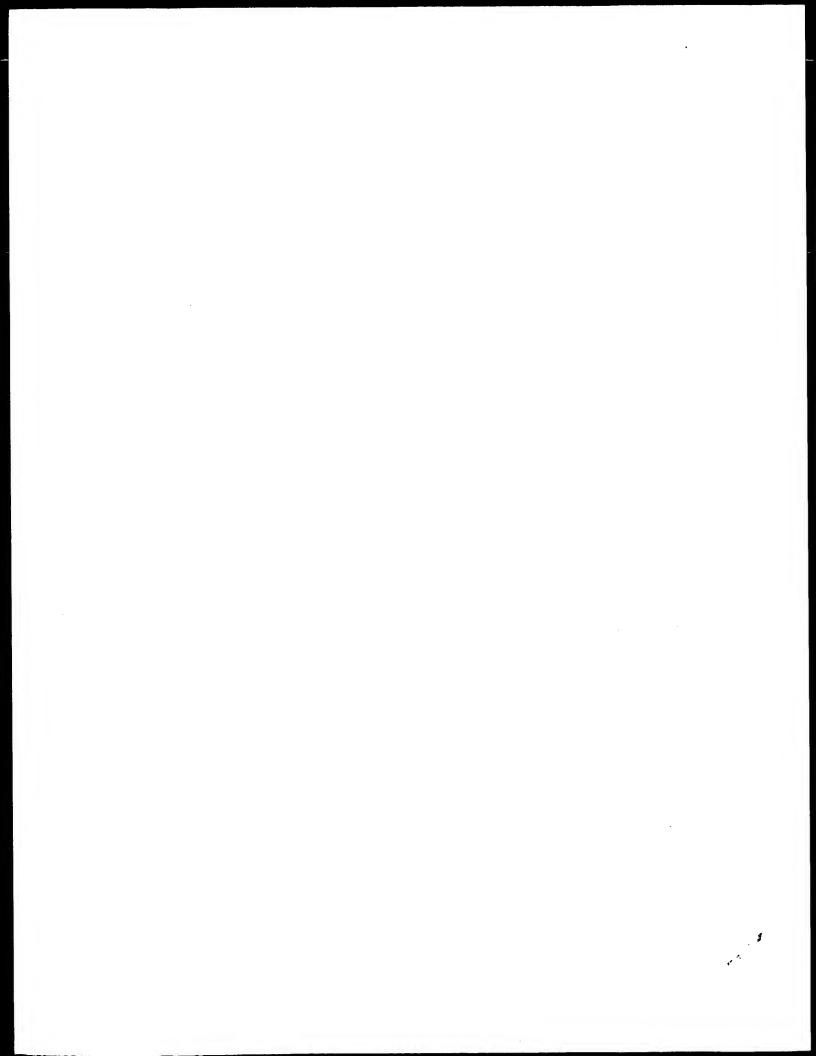
PRIOR FILING DATE: 1998-12-16

NUMBER: OF SEQ ID NOS: 31

SEQ ID NO 6: 31-16

NUMBER: DATE: 1998-12-16

NUMBER: 1998-10
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February 12, 2003, 10:45:03; Search time 11 Seconds (without alignments)
44.130 Million cell updates/sec
GenCore version 5.1.3
(c) 1993 - 2003 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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1 PPLSQETFSDLWKLLPENG 19
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length: 19
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Published Applications AA:*

| 'cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
| 'cgn2_6/ptodata/1/pubpaa/PCT NEW_PUB.pep:*
| 'cgn2_6/ptodata/1/pubpaa/NSO6_NEW_PUB.pep:*
| 'cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
| 'cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
| 'cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
| 'cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
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| 'cgn2_6/ptodata/1/pubcomBCOMBCOMB.pep:*
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RESULT 2 US-09-732-384-7 ; Sequence 7, Application US/09732384 ; Patent No. US20020132977A1

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STRANDEDNESS: single
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US-10-155-059-13
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US-09-732-384-6
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US-09-214-371-1
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LENGTH: 19
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JOSt, Christine
TITLE OF INVENTION: METHODIES OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Ouery Match 87.6%; Score 92; DB 10; Length 19; Best Local Similarity 100.0%; Pred. No. 6.6e-08; Matches 17; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION SATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
GENERAL INFORMATION:
APPLICANT: Yuan, Zhi-Min
APPLICANT: Yuan, Zhi-Min
APPLICANT: Gu, JiJie
ITTLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
FRIOR PILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELECOMMUNICATION INFORMATION: TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Disket
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 27
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COUNTRY: USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REPRENCE: 4-2093/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR FILING DATE: 1997-07-04
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Description of Artificial Sequence:peptide US-09-214-371-1
                                                                                                      Query Match 85.7%; Score 90; DB 12; Length 19; Best Local Similarity 88.9%; Pred. No. 1.3e-07; Matches 16; Conservative 1; Mismatches 1; Indels
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84.8%; Score 89; DB 10; Length 19

Best Local Similarity 94.1%; Pred. No. 1.9e-07;

Matches 16; Conservative 1; Mismatches 0; Indels
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APPLICANT: Yuan, Zhi.Min
APPLICANT: Yuan, Zhi.Min
APPLICANT: Gu, JiJie
IITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21500-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 6
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 1, Application US/09214371B; Patent No. US20010018511A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 6, Application US/09732384; Patent No. US20020132977A1; GENERAL INFORMATION:
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SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                1 PPLSQETFSELWNLLPEN 18
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Patent No. US20010018511A1
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                     GENERAL INFORMATION:
APPLICANT: Lane, D
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LENGTH: 15
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LENGTH: 12
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                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Kaelin, William
JOSt, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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      OTHER INFORMATION: fragment not in inhibitory p53 polypeptide
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70.5%; Score 74; DB 12; Length 16;
Best Local Similarity 81.2%; Pred. No. 2.8e-05;
Matches 13; Conservative 1; Mismatches 2; Indels
                                                                   Length 15;
                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Diskette
COMPUTER: IBM Compactible
COMPUTER: IBM Compactible
COMPUTER: SYSTEM: Windows
SOFTWARE: FastSED for Windows
SOFTWARE: PastSED for Windows
CURRITATION DATA:
PRIOR PAPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
ATTORNEY/AGENT INCRMATION:
                                                              73.3%; Score 77; DB 10; L
100.0%; Pred. No. 9.4e-06;
tive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Bisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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Patent No. US20020147173A1
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
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                                                                                      Best Local Similarity 100.0
Matches 14; Conservative
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                                                                                                                                                     5 QETPSDLWKLLPEN 18
                                                                                                                                                                                            1 OETFSDLWKLLPEN 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
ZIP: 02110
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               ; US-09-732-384-6
                                                                         Query Match
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APPLICANT: Goueli, Said A
TITLE OF INVENTION:
CURRENT APPLICATION Quantitation of Individual Protein Kinase Activity
TITLE OF INVENTION: Quantitation of Individual Protein Kinase Activity
CURRENT APPLICATION NUMBER: US/09/950,692
CURRENT FILING DATE: 2001-06-27
PRIOR FILING DATE: 1994-03-04
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin Ver. 2.0
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                      APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Bascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION INMBER: US/09/214,371B
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR PILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ) OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: polypeptide
US-09-950-692-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       62.9%; Score 66; DB 10; Length 12; 100.0%; Pred. No. 0.00033; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 22, Application US/10155059
; Patent No. US20020147173A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        // Sequence 6, Application US/09950692
// Patent No. US20020106701A1
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
Lane, David
Bottger, Volker
Bottger, Angelica
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Best Local Similarity 100.
Matches 12; Conservative
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                                                 NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE ANTIBODIES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               57.1%; Score 60; DB 12; Length 18; 76.9%; Pred. No. 0.0041; ative 2; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURSENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION - CURNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1999
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Elsensetin, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMUNICATION INFORMATION:
TELEPHONE: 617-345-654
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/214,371B CURRENT FILING DATE: 1999-03-26 PRIOR APPLICATION NUMBER: PCT/EP97/03549 PRIOR FILING DATE: 1997-07-04 NUMBER OF SEQ ID NOS: 83 SOFTWARE: PATENTIN Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE DESCRIPTION: SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
TITLE OF INVENTION: Inhibitors of the I
FILE REFERENCE: 4-20937/A/PCT
                                                                                                                                        ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                 STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 39, Application US/09214371B
Patent No. US20010018511A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 18 amino acids
                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
                                                                                                             CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
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Best Local Similarity 76.99
Matches 10; Conservative
                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                     CITY: Boston
                                                                                                                                                                                                                                     COUNTRY: USA
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Gaps
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APPLICANT: Kim, Seong-Tae
APPLICANT: Kim, Seong-Tae
APPLICANT: Lim, Dae-Sin
APPLICANT: Lim, Dae-Sin
APPLICANT: Lim, Dae-Sin
APPLICANT: Lim, Dae-Sin
APPLICANT: St. Jude Children's Research Hospital
TITLE OF INVENTION: ATM Kinase Modulation for Screening and
TITLE OF INVENTION: Therapies
FILE REPRENCE: 2477/17142
CURRENT FILING DATE: 2001-12-17
PRIOR FILING DATE: 1999-09-21
PRIOR FILING DATE: 1999-09-21
PRIOR FILING DATE: 1999-02-10
NUMBER OF SEQ ID NOS: 49
SOFTWARE: FASTESQ for Windows Version 3.0
                                                                                   FEATURE: OTHER INFORMATION: Description of Artificial Sequence: peptide
                                                                                                                                                                                                                                                                                                             56.2%; Score 59; DB 10; Length 16; 100.0%; Pred. No. 0.0051; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            54.3%; Score 57; DB 9; Length 14; 100.0%; Pred. No. 0.0089; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Picksley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Garcia-Echeverria, Carlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 7, Application US/09214371B; Patent No. US20010018511A1; GENERAL INFORMATION:
APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
                                                                                                                                                     LOCATION: (1)

OTHER INFORMATION: X = Biotin-Ser
NAME/KEY: VARLANT
LOCATION: (16)
US-09-214-371-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 7, Application US/10024123; Publication No. US20030022263A1; GENERAL INFORMATION:
                                        TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100.
Matches 11; Conservative
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Best Local Similarity 100.(
Matches 11, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Kastan, Michael APPLICANT: Canman, Christ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Furet, Pascal
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                                                                                                                                                                                                                                                                                                                                                                                                                                  S QETFSDLWKLL 15
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                                                                                                                                NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-10-024-123-7
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US-09-214-371-7
SEQ ID NO 39
LENGTH: 16
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LENGTH: 14
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                                                                                                                                                                                                                                                                                                                  Query Match
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US-09-214-371-40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Pickeley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Garcia-Ebreveria, Carlos
APPLICANT: Puret, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REPERENCE: 4-20337/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR PILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
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                                                                                                                                                                                                                                                       ) OTHER INFORMATION: Description of Artificial Sequence:peptide US-09-214-371-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Match 51.4%; Score 54; DB 10; Length 12; Local Similarity 100.0%; Pred. No. 0.021; les 10; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                     52.4%; Score 55; DB 10; Length 12; 83.3%; Pred. No. 0.015; tive 0; Mismatches 2; Indels
PILE REFERENCE: 4-20937/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEO ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LOCATION: (1)
OTHER INFORMATION: X = Ac-Gln
NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
                                                                                                                                                                                                   TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 83.3
Matches 10; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                        5 OETFSDLWKLLP 16
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SEQ ID NO 24
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LENGTH: 12
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GENERAL INFORMATION:

APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Hochkeppel, Heinz-Kurt
CURRENT ENTRE OF INVENT: 119901, 4000, 214, 371B
CURRENT APPLICATION NUMBER: US/09/214, 371B
CURRENT FILING DATE: 1999-03-26

CURRENT APPLICATION NUMBER: CAT/EP97/03549
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Stephen
APPLICANT: Bottger, Stephen
APPLICANT: Picksley, Stephen
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REFERENCE: 4-20937/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1997-07-04
PRIOR FILING DATE: 1997-07-04
NUMBER: PatentIn Ver. 2.0
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 41
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence:peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 51.4%; Score 54; DB 10; Length 16; Best Local Similarity 90.9%; Pred. No. 0.029; Matches 10; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 40, Application US/09214371B; Patent No. US20010018511A1
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LOCATION: (1)
OTHER INFORMATION: X = Biotin-Ser
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OTHER INFORMATION: X = Pro-NH2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; LOCATION: (16); COTHER INFORMATION: X = Pro-NH2
US-09-214-371-41
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PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO LENGTH: 16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: VARIANT
LOCATION: (1)
OTHER INFORMATION: Biotin-Ser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: VARIANT LOCATION: (16)
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US-09-214-371-40
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Query Match 50.5%; Score 53; DB 10; Length 16;
Best Local Similarity 90.9%; Pred. No. 0.041;
Matches 10; Conservative 0; Mismatches 1; Indels 0; Gaps 0;
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Oy 5 QETFSDLWKLL 15 | | | | | | | | | | Db 5 QPTFSDLWKLL 15 Search completed: February 12, 2003, 10:50:08 Job time : 12 secs

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p53 protein bindin
Peptide #1. Unide
p53 DR 3a motif bi
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p53 N-terminal pep
p53/RB interaction
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Immunodominant epi
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p53 homologue TIP
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                                                                                                                                 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A_Geneseq_101002:*

| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:*
| SIDS2/gcgdata/geneseqf_embl/AA1981.DAT:*
| SIDS2/gcgdata/geneseqp-embl/AA1981.DAT:*
| SIDS2/gcgdata/geneseqp-embl/AA1981.DAT:*
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                            Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                     908470 segs, 133250620 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                                       OM protein - protein search, using sw model
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AAR89914
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AAW82319
AAW37228
AAW47074
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AAR54909
                                                                                                                                                                                                                                                           105
1 PPLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                                                               BLOSUM62
Gapop 10.0 , Gapext 0.5
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Match Length DB
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seq length: 19
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73.33.3
73.33.3
73.33.3
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Maximum DB
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di.	e #3.		ヹ	#11.	#4	5.	83	ract	Peptide #6. Unide	#8	Ligand #1. Uniden	Peptide #2. Unide	Human p53 wild-typ	Human oncogenic pr		Mdm/hdm antagonist			Human oncogenic pr	Promega protein ki	Tumour suppressor		p53 DR supermotif		Human oncogenic pr	•ط	Mdm/hdm antagonist	Mdm/hdm antagonist	Mdm/hdm antagonist	Human oncogenic pr		Mdm/hdm antagonist	Mdm/hdm antagonist	
AAG89730 AAY45227	5	AAB29163	ABB05528	AAB29167	AAB29160	AAB29161	AAY99001	AAY57799		AAB29164	AAB29149	AAB29158	AAW37181	AAW37188	AAB17075	AAB17087	ABB73170	ABB73182	AAW37201	AAR83012	AAY95859	ABB09246	AAG89447	AAM52269	AAW37190	AAB17077	AAB17089	ABB73172	ABB73184	AAW37203	AAW37189	AAB17076	AAB17088	
22	21	21	23	21	21	21	21	21	21	21	21	21	19	19	21	21	23	53	19	16	21	23	22	23	19	21	21	23	23	19	19	21	21	
15	15	15	14	15	15	15	15	13	15	15	16	16	12	12	12	12	12	12			15													
	70.5						9.19	66.7	65.7											ä	61.9	Η.	φ.	ω.	œ.	œ.	ω,	œ,	œ.	ω,	7.	7.	57.1	
77		74	73	73	71	71	71		69	69	66.5	ė.	99	99	99	99	99	99	99	65	65	65	62	61					61					
11	13	14	15	16	17	18	19	20	21	22	23	24	25	26	27	28	29	30	31	32	33	34	35	36	3.7	38	39	40	41	42	43	44	45	

ALIGNMENTS

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p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.
                                                                                                                                                                                                                                                 New substance with a mdm2 binding domain and coupling partner -useful for stabilising in cells without an efficient mdm2-mediated degradation pathway
                AAW82321 standard; Peptide; 19 AA.
                                                                                                                                                            98WO-GB01140.
                                                                                                                                                                            97GB-0008089.
                                                  (first entry)
                                                                  p53 homologue TIP peptide
                                                                                                                                                                                                                                 WPI; 1998-609975/51.
                                                                                                                                                                                             (UYDU-) UNIV DUNDEE
                                                   22-FEB-1999
                                                                                                                       WO9847919-A1
                                                                                                                                                           20-APR-1998;
                                                                                                                                                                            22-APR-1997;
                                                                                                                                         29-OCT-1998
                                                                                                       Synthetic.
                                  AAW82321;
                                                                                                                                                                                                              Lane DP;
RESULT 1
         AAW8232
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Disclosure; Fig 1; 52pp; English

This sequence is a peptide homologue of a region of p53 which binds to mand. This peptide is used in the construction of a movel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells on or overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 unction. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.

19 AA; Sequence

Gaps ö Length 19; Indels 100.0%; Score 105; DB 19; 100.0%; Pred. No. 5.7e-10; tive 0; Mismatches 0; Local Similarity 100. Query Match Matches

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1 PPLSQETFSDLWKLLPENG 19 1 PPLSÓETFSDLWKLLPENG 19 g ઠે

AAW82319 standard; Peptide; 19 AA. AAW82319

22-FEB-1999 (first entry) AAW82319;

p53 homologue TIP peptide.

p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.

Synthetic.

WO9847525-A1

29-OCT-1998.

98WO-GB01144. 20-APR-1998; 97GB-0008092. 22-APR-1997;

(UYDU-) UNIV DUNDEE.

Lane DP;

WPI; 1998-609932/51.

New agents which inhibit interaction of p53 and mdm2 - useful for activating p53, e.g. for treating cancers, viral conditions or other conditions associated with non functional p53 or mdm2

Disclosure, Fig 1; 52pp; English.

This sequence is a peptide homologue of a region of p53 which binds to mand. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in the preparation of a therapeutic for activating p53 where the population of cells of out overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by reducing the clearance of p53 by mdm2, and can be used to activate p53 unction. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.

Sequence 19 AA;

Query Match

100.0%; Score 105; DB 19; Length 19;

· 0 MDM2; oncogenic protein; p53; human; inhibition; interaction; cancer; tumour; diagnosis; binding; viral infection; Elisa TIP assay. Gaps Garcia-echeverria C; . 0 Indels p53 N-terminal peptide fragment for Elisa TIP assay. Pred. No. 5.7e-10; Furet P, 0; Mismatches (CANC-) CANCER RES CAMPAIGN TECHNOLOGY. Boettger V, Chene P, Lane D, Picksley S; AAW37228 standard; peptide; 18 AA. Best Local Similarity 100.0%; 97WO-EP03549. 97GB-0007041. 96GB-0014197. 1 PPLSQETFSDLWKLLPENG 19 1 PPLSOETFSDLWKLLPENG 19 20-JUL-1998 (first entry) Conservative WPI; 1998-100996/09. (NOVS) NOVARTIS AG. Hochkeppel H, 04-JUL-1997; 07-APR-1997; WO9801467-A2. Homo sapiens 05-JUL-1996; 15-JAN-1998. Boettger A, 19; AAW37228; Matches RESULT 3 ò g

Compounds binding to MDM2 protein and inhibit its interaction with p53 - useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents

Disclosure; Page 34; 45pp; English.

This represents a p53 N-terminal peptide fragment used in an Blisa TIP assay for analysing the interaction between human oncogenic protein MDM2 can diversion provides peptide derivatives capable of binding to the human MDM2. These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated levels of MDM2. The peptides may be used to identify molecules that bind to MDM2 and to identify and design inhibitors of MDM2, p53 binding. They can also be used to purify binding partners especially MDM2, diagnose disease by measuring levels of MDM2 in blood of cancer and leukaemia contexactions, especially tumours and viral infections. The peptides can interactions, especially tumours and viral infection. By interfering with MDM2/p53 interaction, the peptides can activate p53 function and cacumulation in normal cells. The peptides which mimic the MDM2 binding context in p53, have a significantly greater blocking activity compared context in the propersion of the models of the models of the models of the mid-type p53.

18 AA; Sequence

Gaps .; 0 Length 18; 0; Indels 94.3%; Score 99; DB 19; L 100.0%; Pred. No. 4.9e-09; tive 0; Mismatches 0; Local Similarity 100. nes 18; Conservative Query Match Matches

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1 PPLSQETFSDLWKLLPEN 18

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RESULT 4

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(first entry)
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Best Local Similarity 100.
Matches 15; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             mRNA in lymphocytes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        15 AA;
                                                                                                                                                                                 therapy; human.
                                                                                                                                                                                                                                                                                                                                           15-DEC-1997;
                                                                                                                                                                                                              Homo sapiens
                                                                                                                                                                                                                                              WO9931231-A1
                                                                                                                                                                                                                                                                                                              14-DEC-1998;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     29-NOV-1994
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                                06-SEP-1999
                                                                                                                                                                                                                                                                              24-JUN-1999.
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                                                                                                RB18A; p53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        neoplasia.
 AAY06310;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR54909;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                           Frade R;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 6
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   a cell enters apoptosis. Apoptosis is prevented if interaction occurs. The interaction is between regions 1-71 or 290-393 of p53 and region 379-928 of Rb. The invention provides methods to identify compounds able to modulate interaction or binding between p53 and Rb protein. The method and detecting interaction/binding between them. These inhibitory compounds are used in human or veterinary medicine to modulate p53 and c11 ty and processes, specifically for inducing apoptosis in tumour cells (possibly also in cells infected by virus), in vivo or in vitro. Expression of these modulators by gene therapy methods is also contemplated. Other activities that can be affected are transcription repression, G1 arrest, DNA repair, homologous recombination and stabilise p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         a p53 protein and a retinoblastoma (RB) protein. This peptide corresponds to residues 11-29 of human p53 (AAW47079). The interaction between p53 and Rb is found to be critical for determining whether or not
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This peptide fragment of p53 is an inhibitor of the interaction between
                                                                                                                                                                             p53/RB interaction inhibiting peptide 6 (residues 11-29 of human p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Identifying compounds that modulate interaction of p53 and Rb protein - or those that bind to Rb protein, used to induce apoptosis, specifically for treatment of tumours
                                                                                                                                                                                                            Retinoblastoma gene; RB; p53 protein; interaction; inhibitor; tumour; apoptosis; modulator; medicine; veterinary; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   92.4%; Score 97; DB 18; Length 19; 94.4%; Pred. No. 1.1e-08; ive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                         CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAY06310 standard; Peptide; 15 AA.
                                                                             AAW47074 standard; peptide; 19 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 36; Page 63; 83pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                           96GB-0008937
                                                                                                                                                                                                                                                                                                                                                                           97WO-GB01168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2 PPLSQETFSDLWKLIPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 PPLSQETFSDLWKLLPEN 18
1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 1997-549887/50.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     19 AA;
                                                                                                                                                                                                          Retinoblastoma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kouzarides T;
                                                                                                                                            19-MAY-1998
                                                                                                                                                                                                                                                                            Homo sapiens.
                                                                                                                                                                                                                                                                                                            W09741433-A1
                                                                                                                                                                                                                                                                                                                                                                           29-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                           29-APR-1996;
                                                                                                                                                                                                                                                                                                                                           06-NOV-1997
                                                                                                                                                                                                                                                             Synthetic
                                                                                                            AAW47074;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen; transformation-associated protein.
                                                                                                                 wound healing; graff rejection; reperfusion injury;
wyocardial infarction; stroke; traumatic brain injury;
neurodegenerative disease; ischaemia; toxaemia; infection; AIDS;
hepatitis; breast cancer; ovarian cancer; colon cancer; diagnosis;
                                                                           regulatory protein; apoptosis; neoplasia; inflammation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This synthetic peptide corresponds to amino acids 13-27 of human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New p53 regulatory protein (RB18A) useful as, e.g. sources of and primers to detect the transcription rate and abundance of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   77.1%; Score 81; DB 20; Length 15; 100.0%; Pred. No. 3.2e-06;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example 1; Page 25; 87pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR54909 standard; peptide; 15 AA.
Human p53 peptide (aa13-27).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           98WO-EP08560.
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AAY06310
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Matches

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Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New peptides of the invention which interfere with the binding of the p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein binding motif between amino acids 16-33 (AAR89914). The peptides were identified by modifying the p53 consensus binding sequence by substitution of an Alanine at each pos. and identifying which amino acid changes altered binding to expressed MDM2. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and other malignancies
Peptides derived from the N-terminal (amino acids 1-112) or the C-terminal (amino acids 350-393) of protein p53 which specifically acade with anti-p53 antibodies in patients with cancer or precancerous conditions are claimed. The peptides (AARS4907-RS4921) are useful for detecting and monitoring cancerous and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         73.3%; Score 77; DB 17; Length 15; 100.0%; Pred. No. 1.4e-05; tive 0; Mismatches 0; Indels
                                                                                                                                          Length 15;
                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                p53 protein binding motif corresp. to residues 16-30.
                                                                                                                                       73.3%; Score 77; DB 15; I
100.0%; Pred. No. 1.4e-05;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                           AAR89914 standard; peptide; 15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95WO-GB01719.
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                                                                                                                                       Query Match
Best Local Similarity 100.
Matches 14; Conservative
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                                                                                                                                                                                                    QETFSDLWKLLPEN 18
                                                                                                                                                                                                                         1 QETFSDLWKLLPEN 14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Lane DP, Picksley SM;
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es 14; Conserv
                                                                                                           15 AA;
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20-JUL-1994;
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                                                                              conditions.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cancer; pre-cancerous state; detection; diagnosis; human p53 gene; immunodominant epitope; human cellular tumour antigen; transformation-associated protein.
                                                                                                                                                                                                 New immuno:dominant epitope(s) of protein p53 - for detecting and monitoring antibodies indicative of cancer and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      New immuno:dominant epitope(s) of protein p53 - for detecting and monitóring antibodies indicative of cancer and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
                                                                                                                                                                                                                                                                                                                                                                                                                             73.3%; Score 77; DB 15; Length 15;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Immunodominant epitope from p53 N-terminal.
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                                                                                                                                                                                                                                                           Claim 5; Page 42; 62pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 5; Page 42; 62pp; French.
                                                                                                                                       Legros Y, Lubin R, Soussi T;
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                                              93WO-FR01082.
                                                                             92FR-0013110.
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                                                                                                          (EURO-) LAB EUROBIO SA.
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                                                                                                                                                                    WPI; 1994-167463/20
                                                                                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
nes 14; Conserv
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                                              02-NOV-1993;
                                                                           02-NOV-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens
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                   11-MAY-1994
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vaccine; epitope; cytostatic.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Assays and screening methods based on direct interaction between FHA domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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cytotoxic T lymphocyte; T cell; immune response; tumour; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ·.
                                                                                                                                                                                                                         Fork head associated; FHA; domain; transcriptional control;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 15;
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; Pred. No. 1.4e-05;
                                                                                                                                                                                                                                                       DNA replication; DNA repair; cell cycle control
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100.0%; Pred. No. 1...
0; Mismatches
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AAB29157 standard; Peptide; 15 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-MAR-2000; 2000WO-GB01024.
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                                                                                                                 (first entry
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (KUDO-) KUDOS PHARM LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Jackson SP, Durocher D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 PPLSQETFSDLWKL 15
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                                                                                                                                                                                                                                                                                                               Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         28-JUN-1999;
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                                                                                                                 02-FEB-2001
                                                                                                                                                                         Peptide #1.
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                                                     AAB29157;
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the present invention describes isolated preparer 23 epicops (1). Also described are: (1) a cytotoxic T lymphocyte (CTL) that is cultured in vivo and binds to a complex of (1); (2) a peptide (II) comprising (I) and a second epitope and has less than 50 contiguous amino acids, (3) a vaccine composition comprising (II), a unit dose of a peptide with less than 50 contiguous amino acids with 100% identity to the native peptide sequence of p53, and a pharmaceutical excipient; (4) an isolated nucleic acid encoding (I); and (5) an isolated nucleic acid encoding (II) and (5) an isolated nucleic acid encoding (II). (I) has cytostatic activity and can be used in vaccines. The vaccine composition is useful for treating or preventing cancer. (I) and (II) are suseful as diagnostic agents and for evaluating immune responses. Unlike conventional epitopes, immunosuppressive epitopes that may be present in whole antigens can be avoided with the use of the vaccine truther, to modify the composition of the epitopes and turther, to modify the composition of the epitopes and immunogenicity. The possible pathological side effects caused by intrinsic biological activity, are eliminated. Add89363 to AdG89777 represent amino acid sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes isolated prepared P53 epitopes (I). Also
                                                                                                                                                                                                                                                                                                                                                                  Epitope-based vaccines comprising P53 epitope having a specified sequences, useful for treating and preventing cancer, the epitopic peptides is useful as diagnostic agents and for evaluating immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Human; p53; tumour suppressor; human leukocyte antigen; HLA; CTL; cytotoxic T lymphocyte; T cell; immune response; tumour; cancer; vaccine; epitope; cytostatic.
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                                                                                                                                                                                                                                                                             Chesnut R,
                                                                                                                                                                                                                                                                           Fikes J, Sette A, Sidney J, Southwood S,
Keogh E;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 113; 138pp; English.
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                                                                                                                                                                 11-DEC-2000; 2000WO-US33629.
                                                                                                                                                                                                      99US-0458297.
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                                                                                                                                                                                                                                          (EPIM-) EPIMMUNE INC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        present invention.
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                                     Homo sapiens.
                                                                                                                                                                                                      10-DEC-1999;
                                                                                                                             14-JUN-2001.
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                                                       Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                 response
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention describes isolated prepared P53 epitopes (I). Also described are: (1) a cytotoxic T lymphocyte (CTL) that is cultured in vivo and binds to a complex of (I); (2) a peptide (II) comprising (I) and a second epitope and has less than 50 contiguous amino acids; (3) a vaccine composition comprising (II), a unit dose of a peptide with less sequence of p53, and a pharmaceutical excipient; (4) an isolated nucleic acid encoding (I); and (5) an isolated nucleic acid encoding (II). (I) has cytostatic activity and can be used in vaccines. The vaccine composition is useful for treating or preventing cancer. (I) and (II)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          are useful as diagnostic geners and for evaluating immune responses. Unlike conventional epitopes, immunosuppressive epitopes that may be present in whole antigens can be avoided with the use of the vaccine composition of (I). The ability to combine selected epitopes and further, to modify the composition of the epitopes enhances the immunogenicity. The possible pathological side effects caused by infectious agents or whole protein antigens, which might have their own intrinsic biological activity, are eliminated. AAG89363 to AAG89747 represent amino acid sequences used in the exemplification of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                               Epitope-based vaccines comprising P53 epitope having a specified sequences, useful for treating and preventing cancer, the epitopic peptides is useful as diagnostic agents and for evaluating immune
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Immunisation, immunogenicity enhancer; rabies virus strain ERA;
immunódominant T-helper cell determinant; nucleoprotein; antibody;
post-translational modification; p53; cancer; diagnosis.
                                                                                                                                                                                                                                                                                                           Fikes J, Sette A, Sidney J, Southwood S, Chesnut R, Celis E, Keogh B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   73.3%; Score 77; DB 22; Length 15; 100.0%; Pred. No. 1.4e-05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 5; Page 131; 138pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAY45227 standard; peptide; 17 AA.
                                                                                                                                                                11-DEC-2000; 2000WO-US33629.
                                                                                                                                                                                                                 99US-0458297.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  06-JAN-2000 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 PPLSQETFSDLWKL 14
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                                                                                                                                                                                                                                                             (EPIM-) EPIMMUNE INC
                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2001-381493/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   p53 peptide 9-25.
                                                                   WO200141788-A1.
Homo sapiens.
                                                                                                                                                                                                                 10-DEC-1999;
                                                                                                                 14-JUN-2001.
                    Synthetic.
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The present invention describes a polypeptide comprising a portion of a by3 comprising a post-translationally modified amino acid residue where the modification is a phosphorylation, glycosylation or prenylation, and an immunogenicity enhancer. The polypeptide is used to detect anti-p53 autoantibodies in a human patient. The antibody can be used to determine the post-translational modification state of a p53 protein. These are used especially in the diagnosis of cancer. The present sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Assays and screening methods based on direct interaction between FHA domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                               Methods of detecting post-translational modification level of p53, useful for diagnosis of cancer \, -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ٥,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fork head associated, FHA, domain, transcriptional control, DNA replication; DNA repair, cell cycle control.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        73.3%; Score 77; DB 20; Length 17; 100.0%; Pred. No. 1.6e-05; live 0; Mismatches 0; Indels
                                                                                                                                            Thurin M, Hoffman R;
                                                                                                                                                                                                                                                                    Disclosure; Page 45; 59pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAB29159 standard; Peptide; 15 AA.
                                                                    98US-0077512
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            17-MAR-2000; 2000WO-GB01024.
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99GB-0015075.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity 100.
es 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (KUDO-) KUDOS PHARM LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 PPLSQETFSDLWKL 17
                                                                                                                                                                               WPI; 1999-561693/47.
                                                                                                        (WIST-) WISTAR INST.
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                                                                                                                                         Ertl H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      17 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WO200057184-A2.
                                11-MAR-1999;
                                                                      11-MAR-1998;
16-SEP-1999.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  invention.
                                                                                                                                           Otvos L,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB29159;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
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15 AA;

Sequence

X S

Gaps . 0

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The present invention relates to assays and screening methods based on a direct interaction between fork head associated (FHA) domains and phosphorylated polypeptides, for characterizing the binding of these molecules. FHA peptides may be useful for treating medical conditions associated with defects in transcriptional control, DNA replication, DNA repair, cell cycle control or other cellular process. The method may provide valuable insights into checkpoint signalling, has important implications for the functions of other FHA domain-containing proteins and provides basis for new lines of therapy. The present sequence is a peptide used in the present invention.
                                                           The present invention relates to assays and screening methods based on a direct interaction between fork head associated (FHA) domains and phosphorylated polypeptides, for characterizing the binding of these molecules. PHA peptides may be useful for treating medical conditions associated with defects in transcriptional control, DNA replication, DNA repair, cell cycle control or other cellular process. The method may provide valuable insights into checkpoint signalling, has important implications for the functions of other FHA domain-containing proteins and provides basis for new lines of therapy. The present sequence is a peptide used in the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Assays and screening methods based on direct interaction between FHA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fork head associated, FHA, domain, transcriptional control, DNA replication; DNA repair; cell cycle control.
                                                                                                                                                                                                                                                                                                                70.5%; Score 74; DB 21; Length 15; 92.9%; Pred. No. 4.2e-05; ive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB29163 standard; Peptide; 15 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 2; 92pp; English.
                                 Disclosure, Fig 2; 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17-MAR-2000; 2000WO-GB01024.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     domains and phosphopeptides,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              99GB-0006432.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                    Local Similarity 92.5
es 13; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Durocher D;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (KUDO-) KUDOS PHARM LTD
                                                                                                                                                                                                                                                                                                                                                                                               1 PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                                                                                                                   2 PPLAQETFSDLWKL 15
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            02-FEB-2001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptide #7.

    binding

                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB29163;
                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 14
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                                                                                                                                                                                                                                                                                                                 Ubiquitin dependent proteolysis modulation; cdc4 phospho design motif; CDP motif; cytostatic; nootropic; antiproliferative; cell proliferation; growth; differentiation; cancer; neurodegenerative disorder;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      that targets molecules for ubsquirin dependent processives. (C) have cytostatic, nootropic and antiproliferative activity, Also described is a method for the treatment of a disease or condition where affected is cells have a defective protein, comprising administering (C) to promote degradation of the target protein in cells by ubiquirin dependent proteolysis. (C) can also be used for modulating the proliferation, growth and/or differentiation of cells. (C) can be used to modulating ubsquirin dependent proteolysis or cell proliferation, growth and or differentiation of cells. (C) can be used to modulate ubiquirin of cells. (C) is useful in the treatment of cancers and neurodegenerative disorders as well as spinal degeneration. The present sequence represents a biotinylated peptide which is used in an example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention describes a cdc4 phospho design (CPD) motif, (C),
                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New Cdc4 Phospho Design motif that targets molecules for ubiquitin dependent proteolysis, is useful for the modulation of cell
                                 .
0
Score 74; DB 21; Length 15;
Pred. No. 4.2e-05;
1; Mismatches 0; Indels
                                  1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Tyers M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "phospho-Thr"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           proliferation i.e. cancer treatment
                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 2; Page 47; 83pp; English.
                                                                                                                                                                                     ABB05528 standard; Peptide; 14 AA.
                                                                                                                                                                                                                                                                                   Biotinylated peptide SEQ ID NO:13,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (MOUN ) MOUNT SINAI HOSPITAL.
70.5%;
92.9%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      04-MAY-2000; 2000US-202166P.
24-JAN-2001; 2001US-263774P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        04-MAY-2001; 2001WO-CA00632.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 from the present invention.
                                                                                                                                                                                                                                                     (first entry)
                                    13; Conservative
                                                                                                 15
                                                                    1 PPLSQETFSDLWKL 14
                                                                                                                                                                                                                                                                                                                                                                     spinal degeneration.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2002-164074/21.
                                                                                               2 PPLSQETFADLWKL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pawson T,
   Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                      ABB05528;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nash P,
                                      Matches
                                                                                                                                                      RESULT 15
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Gaps

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0; Indels

69.5%; Score 73; DB 23; L 100.0%; Pred. No. 5.6e-05; tive 0; Mismatches 0;

Best Local Similarity 100. Matches 13; Conservative

Query Match

Length 14;

- 1 PPLSQETFSDLWK 13 ||||||||||||||| 2 PPLSQETFSDLWK 14 Q Q

Search completed: February 12, 2003, 10:43:48 Job time : 35 secs

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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 10:42:32 ; Search time 28 Seconds Run on:

(without alignments)
139.818 Million cell updates/sec

US-09-403-440A-2

105 1 PPLSQETFSDLWKLLPENG 19 Perfect score: Sequence:

BLOSUM62 Scoring table:

Gapop 10.0 , Gapext 0.5

671580 seqs, 206047115 residues Searched:

Total number of hits satisfying chosen parameters:

5270

Minimum DB seq length: 0 Maximum DB seq length: 19

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

sp_archea:* sp_bacteria:* SPTREMBL 21:*

sp_plant:*
sp_rodent:*
sp_virus:*
sp_vertebrate:*
sp_unclassified:* sp_fungi:*
sp_human:*
sp_invertebrate:*
sp_mammal:*
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sp_norganelle:* sp_rvirus:* sp_bacteriap:* sp_archeap:* 10:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

			de			COLLEGATES	
Result	11t		Query				
2	No.	Score	Match	Match Length DB	DB	ID	Description
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12 7 077919 14 8 09M03 15 13 09FS10 16 12 083967 16 12 084055 17 11 096VS6	27.78	55 4 4 6 10 6 10 6 10 6 10 6 10 6 10 6 10	4 1 4 1 2 2 1 4 1 4 1 4 1	14 2 Q9R518 17 15 Q72017 18 13 013167 19 6 019107 7 8 Q95945 9 4 Q9H3Y3 13 11 Q91XP1
444444	33 21. 3 21. 21.	33 221. 3 221. 3 21.	3 21. 3 21. 2 21. 2 21.	22 21.0 22 21.0 22 21.0 21.5 20.5 21 20.0 21 20.0
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ALIGNMENTS

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Gonzalez., Mendoza G., Ramos F., Garcia G., Moran P., Valadez A.,
Zaragoza M.E., Melendro E.I., Ximenez C.;
Zaragoza M.E., Melendro E.I., Ximenez C.;
"Calreticulin-like molecule in trophozoites of Histolytica HM1-IMMS.";
"Calreticulin-like molecule in trophozoites bank.
"Calle Edward County and Edward Brown and Edward Brown and Edward Brown and Edward Brown Experience of Francisco III BROWELOWENCE ETICULUM LUMEN.

-1- SUBCELLULAR LOCATION: BIDOPLASMIC RETICULUM LUMEN.
-1- SUBCLOPMENTAL STAGE: IN TROPHOZOITES.
-1- SIMILARITY: BELONGS TO THE CALRETICULIN FAMILY.

RESPECT PROGUEDS CALRETICULIN 1; PRATIAL.

RROSITE; PSOO803; CALRETICULIN 2; PARTIAL.

RROSITE; PSOO805; CALRETICULIN REPEAT; PARTIAL.

RROSITE; PROSOSS; CALRETICULIN REPEAT; PARTIAL.

RROPLASMIC RELICULUM: Calcium-binding.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE 19 AA; 2488 MW; FC90BCAEFE1BA764 CRC64;
                                                            P83003;
01-0CT-2001 (TrEMBLrel. 18, Created)
01-0CT-2001 (TrEMBLrel. 18, Last sequence update)
01-0CT-2001 (TrEMBLrel. 18, Last annotation update)
                                             19 AA
                                                                                                                                             Calreticulin-like protein (Fragment)
Entamoeba histolytica.
                                          PRT;
                                                                                                                                                                                      Eukaryota; Entamoebidae; Entamoeba.
NCBI_TaxID=5759;
                                          PRELIMINARY;
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RESULT 1
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Jiang H., Lucy M.C.;
"Identification of 5' untranslated variants of bovine growth hormone
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                                                                                                                                                                                                                     Larkin D., Serov O., Zhdanova N.; "Mapping of five genes from human chromosome 17 to chromosome hn of the common shrew (Sorex araneus) "; the common shrew (Sorex araneus) "; Acta Theriol. (Warsz) 45:143-146(2000).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=BRBED HOLSTEIN;
BRDLINE=99175163; PubMed=10075683;
Jiang H., Okamura C.S., Lucy M.C.;
"Isolation and characterization of a novel promoter for the bovine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
MCBI_TaxID=9913;
                                                                                                                  Sorex araneus (Eurasian common shrew) (European shrew).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Insectivora, Soricidae, Soricinae, Sorex.
                                                                                                                                                                                                                                                                                                                                                                                                              ;
                                                                                                                                                                                                                                                                                                                                                                        . Match 29.5%; Score 31; DB 6; Length 16; Local Similarity 60.0%; Pred. No. 3.4e+02; les 6; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Jiang H., Lucy M.C.;
"Bovine GH receptor 5' UTR variants.";
Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; AF036297; AAC33315.2; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Submitted (DEC-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Submitted (DEC-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                           SEQUENCE 16 AA; 1951 MW; 775186E3FE5F52E2 CRC64;
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01-MAR-2001 (TrEMBLrel. 16, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                              01-UUN-2001 (TrEMBLrel. 17, Created)
01-UUN-2001 (TrEMBLrel. 17, Last sequence update)
01-UUN-2001 (TrEMBLrel. 17, Last annotation update)
Thyroid hormone receptor alpha (Fragment).
16 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    growth hormone receptor gene.";
J. Biol. Chem. 274:7893-7900(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Growth hormone receptor (Fragment).
PRT;
                                                                                                                                                                                                                                                                                          EMBL; AF314827; AAK13419.1; -.
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 PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bos taurus (Bovine).
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                                                                                                                                                                      NCBI_TaxID=42254;
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MEDLINE=99297341; PubMed=10368952; Campbell N.J.H., Barker S.C.; Campbell N.J.H., Barker S.C.; The novel mitochondrial gene arrangement of the cattle tick, Boophilus microplus: fivefold tandem repetition of a coding region."; MNOL Biol. Evol. 16:732-740(1999).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Parasitiformes; Ixodida; Ixodidae; Boophilus.
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                                                                                                                                                                                                                                                                                                                                                                                         Length 14;
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                                                                                                                                                                                                                                                                                                         14 14 14 14 14 14 AA; 5A45D458FF3A9E27 CRC64;
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Last annotation update)
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
Cytochrome oxidase subunit 2 (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                         Score 30; DB 6; I
Pred. No. 4.2e+02;
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01-MAY-2000 (TrEMBLrel. 13, Last seque
01-UNN-2001 (TrEMBLrel. 17, Last annoi
FAS antigen (CD95 antigen) (Fragment)
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EMBL, AF085281, AAD22521.1; --
EMBL, AF036290, AAC33308.1; --
EMBL, AF036291, AAC33309.1; --
EMBL, AF036291, AAC33310.1; --
EMBL, AF036293, AAC33311.1; --
EMBL, AF036294, AAC33312.1; --
EMBL, AF036296, AAC33313.1; --
EMBL, AF036296, AAC33314.1; --
EMBL, AF036296, AAC33314.1; --
EMBL, AF036396, AAC33314.1; --
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Best Local Similarity 75.v.,
6; Conservative
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SEQUENCE
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STRAIN=CV. DE GRACE; TISSUE=LRAF;
MEDLINE=20181728; PubMed=10715320;
A Peliter J.-B., Friso G., Kalume D.E., Roepstorff P., Nilsson F.,
A Adamska I., van Wijk K.J.;
I "Proteonics of the chloroplast: systematic identification and
targeting analysis of lumenal and peripheral thylakoid proteins.";
I "Proteophara Location" CHLOROPLAST THYLAKOID MEMBRANE LUMEN.
-I- BUSELLULAR LOCATION: CHLOROPLAST THYLAKOID MEMBRANE LUMEN.
-I- DEVELOPMENTAL STAGE: UNFOLDED AND FULLY DEVELOPED LEAVES.
-I- MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN
CHORDLAST: Thylakoid membrane.

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                                                   "Transcription stimulation of the Fas-encoding gene by nuclear factor for interleukin-6 expression upon influenza virus infection."; J. Biol. Chem. 270:18007-18012(1995).
                                                                                                                                                                                                              Muschen M., Re D., Jungnickel B., Diehl V., Rajewsky K., Kuppers R.; "Somatic mutations of the CD95 gene in human B cells as a side-effect
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-JUN-2000 (TrEMBLrel. 14, Created)
01-JUN-2000 (TrEMBLrel. 14, Last sequence update)
01-JUN-2000 (TrEMBLrel. 15, Last annotation update)
Unknown protein from 2D-gage of thylakoid lumen (SpCTIO7) (Fragment).
Pisum sativum (Garden pea).
Bukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta; Spermatophyta; Magnoliophyta; eudicotyledons; core eudicots; Rosidae; eurosids 1; Fabales; Fabaceae; Papilionoideae; Vicieae; Pisum.
NCBL TaxID=3888;
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                                                                                                     SEQUENCE FROM N.A.
Muschen M., Re D., Brauninger A., Wolf J., Hansmann M.L., Diehl V.,
Kuppers R., Rajewsky K.;
"Sometic mutations of the CD95 gene in Hodgkin- and Reed-Sternberg
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           MEDLINE-95355401; PubMed-7543095;
Wada N., Matsumura M., Ohba Y., Kobayashi N., Takizawa T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 26; DB 10; Length 14; Pred. No. 1.9e+03; 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                            25.7%; Score 27; DB 4; Length 11; 66.7%; Pred. No. 1e+03;
                                                                                                                                                                                                                                                                                                                                                                                         1; Indels
                                                                                                                                                                                                                                    of the germinal center reaction.";
Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases.
EMBL; D31968; BAA20850.1;
EMBL; AA279012; CAC35539.1;
EMBL; AA279012; CAC35540.1;
EMBL; AJ279013; CAC35540.1;
                                                                                                                                                                       Submitted (SEP-2000) to the EMBL/GenBank/DDBJ databases
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                                                                                                                                                                                                                                                                                                                                                                                         4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NAR-2002 (TrEMBLrel. 20, Last annotation update)
11-NAR-2002 (TremBlrel. 20, Last annotation update)
11-NAR-2004 (H212), non-structural protein (Seg 8), cooh terminus of nsl (Fragment).
                                                                                                                                                                    Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Sandulache R., Pretsch W., Chatterjee B., Gimbel W., Graw J.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 17;
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"Human oviduct glycoprotein promoter sequence.";
Submitted (FEB-2000) to the EMBL/GenBank/DDBJ databases.
EMBL, AF189710; AAF01065.1; -.
NON TER
SEQÜENCE 18 AA; 2201 MW; 5E8FD91EA210E516 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                           mouse.";
Mamm. Genome 5:777-780(1994).
REDIENCE 17 AA; 1982 MW; E941E0A3F2477D45 CRC64;
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01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                          Last sequence update)
Last annotation update)
17 AA.
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                                           01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last seq
01-DBC-2001 (TrEMBLrel. 19, Last ann
Lactate dehydrogenase-A (Fragment).
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Best Local Similarity 30.8%
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12 WKLLPENG 19
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Bedzyk L.A., Escudero K.W., Gill R.E., Griffin K.J., Frerman F.E.;
"Cloning, sequencing, and expression of the genes encoding subunits of
Paracoccus denitrificans electron transfer flavoprotein.";
J. Biol. Chem. 268:20211-20217(1993).
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MEDLINE=21101849; PubMed=11160087;
Fales L., Kryszak L., Zeilerra-Ryalls J.;
"Control of hema Expression in Rhodobacter sphaeroides 2.4.1: Effect of a Transposon Insertion in the hbdA Gene.";
J. Bacteriol. 183:1568-1576(2001).
                                                                                                                                                                                                     Gaps
                                                                    Parvin J.D., Young J.F., Palese P., "nonsense mutations affecting the lengths of the ns1 nonstructural proteins of influenza a virus isolates."; Virology 128:512-517(1983).
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Bacteria; Proteobacteria; alpha subdivision; Rhodobacter group;
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Pred. No. 3.2e+03;
1; Mismatches 3; Indels
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Influenzavirus A.
Viruses; ssRNA negative-strand viruses; Orthomyxoviridae;
Influenza A viruses.
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                                                                                                                                                          SEQUENCE 16 AA; 1811 MW; 23F434B5AEEFAD93 CRC64;
                                                                                                                                                                                                                                                                                                                              01-WAR-2001 (TrEMBLrel. 16, Last sequence update) 01-DEC-2001 (TrEMBLrel. 19, Last annotation update) Electron transfer flavoprotein subunit A (Fragment)
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01-OCT-2000 (TrEMBLrel. 15, Created)
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                                                    SEQUENCE FROM N.A.
MEDLINE=83303830; Pubmed=6612993;
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                                                                                                          EMBL; K00962; AAA43516.1; -.
InterPro; IPR000256; Flu NS1.
Pfam; PF00600; Flu NS1; 1.
NON TER
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                                NCBI_TaxID=11320;
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01-MAR-2001
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 20, Last annotation update)
101-NAR-2002 (TrEMBLrel. 20, Last annotation update)
10f1uenza A/Swine/Wisconsin/OM/2 (HINI), non-structural protein (Seg 8), COOH terminus of NSI (Fragment).
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Inonesnes mutations affecting the lengths of the nsl nonstructural proteins of influenza a virus isolates.";
Virology 128:512-517(1983).
Wirology 128:512-517(1983).
InterPro; IPR00255; Flu Nsl.
Pfam; PF00600; Flu Nsl.
                                                                                    Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
Mitochondrial 28S ribosomal protein S5 (MRP-S5) (Fragments).
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                                                                                                                                                                                                          SEQUENCE, AND MASS SPECTROMETRY.
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Similarity 57.1%;
4; Conservative
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NCBI_TaxID=11320;
                                                                        Bos taurus (Bovine).
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Influenzavirus A.
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Influenza A/berkeley/1/68 (H2n2), non-structural protein (Seg 8), cooh terminus of ns1 (Fragment).
Influenzavirus A.
Viruses, ssRNA negative-strand viruses; Orthomyxoviridae;
Influenza A viruses.
                                                                                                                                                                                          Dioscorea tenuipes.
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Dioscoreales, Dioscoreaceae,
                                                                                                                                                                                                                                                                                                                                                                                                                 Terauchi R., Terachi T., Miyashita N.;
"DNA polymorphism aut the Pgi locus of a wild yam, Dioscorea tokoro.";
Genetics 147:1899-1914 (1997).
EMBL; AB006004; BAA23204.1; -.
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Parvin J.D., Young J.F., Palese P.;
Parvin J.D., Young J.F., Palese P.;
Proteins of influenza a virus isolates.";
Virology 128:512-517(1983).
EMBL; K00957; AAA43518.1; -.
InterPro; IPR000256; Flu NS1.
Pfam; PF00600; Flu_NS1; I.
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Pred. No. 3.8e+03;
2; Mismatches 7; Indels
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NON TER 19 19
SEQUENCE 19 AA; 2245 MW; 333500DFC9EDA5F8 CRC64;
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                                                                  01-JAN'1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Phosphoglucose isomerase (EC 5.3.1.9) (Fragment).
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MEDLINE=98072503; PubMed=9409845;
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30.8%;
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Matches 4, Conservative
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NCBI_TaxID=64476;
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Gaps
                                                                                                                                                                                                                                                                                                     MEDLINE=0150243; PubMed=10684923; Saguez C., Lecellier G., Koll F.; Saguez C., Lecellier G., Koll F.; Intronic GIY-YIG endonuclease gene in the mitochondrial genome of Padospora curvicolla: evidence for mobility."; Nucleic Acids Res. 28:1299-1306 (2000). EMBL; Ad349984; CAB7247.1; InterPro; IPR000179; Cyt. b.b6. PROSITE; PS00193; CYT. b.b6. DRINNOWN_1.
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0
                                                                                                                                                            Eukaryota; Pungi; Ascomycota; Pezizomycotina; Sordariomycetes; Sordariales; Lasiosphaeriaceae; Podospora.
NCBL_TaxID=48157;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE 10 AA; 1200 MW; C31A223B437B0772 CRC64;
01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-UN-2001 (TrEMBLrel. 17, Last annotation update)
Cytochrome b (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 80.0
Matches 4; Conservative
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                                                                                                                         Podospora curvicolla.
                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                  Mitochondrion.
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 10:36:28; Search time 29 Seconds (without alignments) 27:174 Million cell updates/sec Run on:

Title: Perfect score: Sequence:

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

112892 seqs, 41476328 residues Searched:

1015 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 19

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_40:* Database : Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

CTIMMADIEC

		de			SUMMARIES	
Result		Query				
No.	Score	Match	Length	DB	ID	Description
-	24	22.9	,	-	HTF HELZE	P16353 heliothis z
7	23	21.9	1	П	AKHX LOCMI	6 locusta mi
m	22	21.0	7	Н	NP1 LYMST	8 lymnaea
4	22	21.0	13	7	NP2_LYMST	lymnaea
S	22	21.0	13	٦	NP4_LYMST	lymnaea
9	22	21.0	13	٦	NP5_LYMST	lymnaea
7	22	21.0	16	٦	IBP4 PIG	sus scro
80	21.5	20.5	12	H	UR2 SCYCA	SCV]
σ	21	20.0	8	Н	HTF2 PERAM	
10	21	20.0	10	-	HTF2_CARMO	
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13	21	20.0	18	7	CPAX BOVIN	
14	20	19.0	80	Н	AKH TABAT	
15	20	19.0	œ	Н	HTF1 PERAM	
16	20	19.0	10	Н	HTF_TABAT	P14596 tabanus atr
	20	19.0	13	1	NP3_LYMST	P80180 lymnaea sta
18	50	19.0	17	-	PH4_PERAM	
19	20	19.0	18	-	PCG6_PACGO	
20	70	19.0	19	Н	AL22_HORSE	
21	20	19.0	19	П	IRBP_CAVPO	
22	20	19.0	19	-	PCG7_PACGO	P82420 pachycondyl
23	19	18.1	80	-1	AKHG_GRYBI	
24	19	18.1	80	1	CLP_THICU	
52	19	18.1	80	-	HTF_TENMO	P25419 tenebrio mo
56	19	18.1	10	-	HTF_NAUCI	P10939 nauphoeta c
27	19	18.1	10	-	URA7 HUMAN	
28	19	18.1	15	Н	- 1	P21225 pisum sativ
29	19	18.1	15	-	- 1	
30	19	18.1	15	~1	PC20_BRANA	9
31	13	18.1	16	Н.	·)	Ф
32	13	18.1	19	Н.		14440
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.04 pinus pinas 83 pinus pinas	57 physcomitre 18 libellula a	25 bandalus bo 25 bacillus th 73 pseudomonas	, 0 0,	40 zea mays (m 27 streptococc	
P81104	P806 P254	F08939 P49325 P55173	P80464 P41493	P80640 P58827	P22583
FKB7_PINPS HS11_PINPS	RBS_PHYPA AKH_LIBAU PDCW_DANBO	SLAP_BACTG POOC_PSEFL	Q2OA_COMTE	UC34 MAIZE ARCA_STRP5	LEC3_PSOSC
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15	12 12 13 13 13 13 13 13 13 13 13 13 13 13 13	10	11	15	15
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ALIGNMENTS

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NP4 LYMST
P80181;
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Lymna-DF-anide 2.
Lymnae ateganalis (Great pond snail).
Lymnaea stagnalis (Great pond snail).
Lymnaeidae; Lymnaea.
NCBL TaxID=6523;
(11)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bukaryota; Metazoa; Mollusca, Gastropoda; Pulmonata; Basommatophora;
Lymnaeidae; Lymnaea.
NCBI_TaxID=6523;
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Johnsen A.H., Rehfeld J.F.;
Jimmangramides, a new family of neuropeptides from the pond snail,
Lymmangramides. a new family of neuropeptides from the pond snail,
invertebrates?";
                           Eur. J. Biochem. 213:875-879(1993).
-!- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
IR; 532471; 532471.
Neuropeptide; Amidation.
MOD_RES 13 13 AMIDATION.
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                                                                                                                                                                                                                   PYRROLIDONE CARBOXYLIC ACID.

        MOD RES
        13
        AMIDATION.

        SEQÜENCE
        13 AA, 1519 MW; 9CA07BA3F5D5B455 CRC64;

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01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
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01-JUL-1993 (Rel. 26, Last sequence update)
01-JUL-1993 (Rel. 26, Last annotation update)
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P80179;
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P80178;
                         SEQUENCE.
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Matches
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NP1_LYMST
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Lymnaea stagnalis (Great pond snail).
Eukaryota; Metazoa; Mollusca; Gastropoda; Pulmonata; Basommatophora;
Lymnaeidae; Lymnaea.
NCBI_TaxID=6523;
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"LymnaDFamides, a new family of neuropeptides from the pond snail,
Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
TISSUB=93238777; PubMed=8477756; Johnsen A.H., Rehfeld J.F.; Johnsen A.H., Rehfeld J.F.; Johnsen A.H., Rehfeld J.F.; LymanDFrantdes, a new family of neuropeptides from the pond snail, Lymaneas stagnalis. Clue to cholecystokinin immunoreactivity in invertebrates?";
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Eur. J. Biochem. 213:875-879(1993).
-!- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
                                                                                                                                                      Length 13;
                                                                                                                                                                                                                                                                                                                                     21.0%; Score 22; DB 1; Length 13; 50.0%; Pred. No. 1.40+03; tive 1; Mismatches 3; Indels
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SEQÜENCE 13 AA; 1503 MW; 9CA07BBB56D5B455 CRC64;
                                                                                                                                                                                                                                                                                              SEQUENCE 13 AA; 1492 MW; 9CA07BA3F5D5A5A5 CRC64;
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01-UUL-1993 (Rel. 26, Last sequence update)
01-UUL-1993 (Rel. 26, Last annotation update)
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MEDLINE=93238777; PubMed=8477756;
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MOD RES 13 13
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Matches 4; Conservative
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HTF2 PERAM
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HTF2_PERAM
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                                                   Johnsen A.H., Rehfeld J.F., "LymnaDFamides, a new family of neuropeptides from the pond snail, Lymnaea stagnalis. Clue to cholecystokinin immunoreactivity in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=92109718; PubMed=1722398;
Coleman M.B., Pan Y.-C.E., Etherton T.D.;
Identification and MI2-terminal amino acid sequence of three insulin-like growth factor-binding proteins in porcine serum.";
Blochem. Blophys. Res. Commun. 181:1131-1136 (1991).
-! FUNCTION: IGFS-BINDING PROTEINS PROLONG THE HALF-LIFE OF THE IG AND HAVE BEEN SHOWN TO EITHER INHIBIT OR STIMULATE THE GROWTH PROMOTING EFFECTS OF THE IGFS ON CELL CULTURE. THEY ALTER THE INTERACTION OF IGFS WITH THEIR CELL SURFACE RECEPTORS.
-! SUBCELLULAR LOCATION: Secreted.
-! SIMILARITY: CONTAINNS 1 THYROGLOBULIN TYPE-I DOMAIN.
-! SIMILARITY: EBLONGS TO THE INSULIN-LIKE GROWTH FACTOR BINDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sus scrofa (Pig).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Cetartiodactyla; Suina; Suidae; Sus.
NCBI_TaxID=9823;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MR-1992 (Rel. 21, Created)
01-MR-1992 (Rel. 21, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
Insulin-like growth factor binding protein 4 (IGFBP-4) (IBP-4)
(IGF-binding protein 4) (Fragment).
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                                                                                                                                                                Eur. J. Biochem. 213:875-879(1993).
-1- SIMILARITY: RELATED TO THE CHOLECYSTOKININ (CCK) FAMILY.
PIR; S32475; S32475.
Neuropeptide, Amidation.
MOD RES
13 13 AMIDATION.
SEQÜENCE 13 AA; 1476 MW; 9CA07BBB56D5A5A5 CRC64;
                                                                                                                                                                                                                                                                                                                                                            21.0%; Score 22; DB 1; Length 13; 50.0%; Pred. No. 1.40+03; vative 1; Mismatches 3; Indels
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Pred. No. 1.8e+03;
1; Mismatches 1; Indels
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9CA07BBB56D5A5A5 CRC64;
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INCETPO: IPR010867; Insl_gro_fac_pr.
INCETPO: IPR010716; Thyraglobulin_1.
PROSITE; PS00222; IGF BINDING; PARTIAL.
PROSITE; PS00484; THYROGLOBULIN_1; PARTIAL.
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#ISSUE=Ganglion;
MEDLINE=93238777; PubMed=8477756;
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13 AA; 1476 MW;
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1es 4; Conservative
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                                                                                                                                         invertebrates?";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 LSOETFSD 10
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ID IBP4_PIG
AC P24854;
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12 AA.

STANDARD;

UR2_SCYCA P35490;

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MEDLINE-84298179; PubMed-6591205;
Scarborough R.M., Jamieson G.C., Kalish F., Kramer S.J., McEnroe G.A.,
Miller C.A., Schooley D.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Blatta orientalis (Oriental cockroach).

Bukaryota, Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
Ilnsecta; Pterygota; Neoptera; Orthopteroidea; Dictyoptera; Blattoidea; Blattidae; Parijaneta.

NCBI_TaxID=6978, 7539, 6976;
                                                                                                                                                                                                                                                                                        MEDLINE=92319231; PubMed=1620290;
Conlon J.M., O'Harte F., Smith D.D., Balment R.J., Hazon N.;
"Purification and characterization of urotensin II and parvalbumin from an elasmobranch fish, Scyliorninus canicula (common dogfish).";
Neuroendocrinology 55:230-235(1992).
-!- FUNCTION: HAS A SUGGESTED ROLE IN OSMOREGULATION AND AS A CORTICOTROPIN-RELEASING FACTOR. PROBABLY INVOLVED IN SMOOTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-1987 (Rel. 26, Last sequence update)
01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
Hypertrehalosaemic factor II (Neuropeptide M-II) (Periplanetin CC-2)
Periplaneta americana (American cockroach),
Leptinotarsa decemiineata (Colorado potato beetle), and
                                                                                               Scyliorhinus canicula (Spotted dogfish) (Spotted catshark).
Wataryota: Metazoa; Chordata; Craniata: Vertebrata; Chondrichthyes;
Elasmobranchii; Galeoncrphii; Galeoidea; Carcharhiniformes;
Scyliorhinidae; Scyliorhinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Structures of two cockroach neuropeptides assigned by fast atom bombardment mass spectrometry."; Biochem. Biophys. Res. Commun. 124:350-358(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SPECIES=P.amerioana;
MEDLINE=85046530; PubMed=6548628;
Witten J.L., Schaffer M.H., O'Shea M., Cook J.C., Hemling M.E.,
Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    "Isolation and primary structure of two peptides with cardioacceleratory and hyperglycemic activity from the corpora cardiaca of Periplaneta americana."

Proc. Natl. Acad. Sci. U.S.A. 81:575-5579 (1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    h 20.5%; Score 21.5; DB 1; Length 12; Similarity 71.4%; Pred. No. 1.6e+03; 5; Conservative 0; Mismatches 1; Indels
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12 Aa; 1526 MW; 804729F9D579CEBA CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                          MUSCLE STIMULATION.
-!- SIMILARITY: BELONGS TO THE UROTENSIN 2 FAMILY.
InterPro; IPR001483; Urotensin II.
PRO2083; Urotensin II; II.
PROSITE; PS00984; UROTENSIN_II; I.
01-JUN-1994 (Rel. 29, Created)
01-JUN-1994 (Rel. 29, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Urotensin II (U-II) (UII)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
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                                                                                                                                                                                                                                                                      TISSUE=Spinal cord;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Best Local Similarity
                                                                                                                                                                                                NCBI TaxID=7830;
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                                                                                                                                                                                                                                         MEDLINE=90253659; PubMed=2340112;
Gaede G., Rinehart K.L. Jr.;
"Primary structures of hypertrehalosaemic neuropeptides isolated from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Extatosoma tiaratum (Stick insect).

Eukaryota, Metazoa, Arthropoda, Mandibulata, Pancrustacea, Hexapoda,
Insecta, Pterygota, Neoptera, Orthopteroidea, Phasmatodea,
Hetronemidae, Carausius.
                                                                                                                                                                                                                                                                                                        the corpora cardiaca of the cockroaches Leucophaea maderae, Gromphadorhina portentosa, Blattella germanica and Blatta orientalis and of the stick insect Extatosoma tiaratum assigned by tandem fast atom bombardment mass spectrometry.";

Biol. Chem. Hoppe-Seyler 371:345-354(1990).

-!- FUNCTION: HYPERTREHALOSAEMIC FACTORS ARE NEUROPEPTIDES THAT ELEVATE THE LEVEL OF TREHALOSE IN THE HEMOLYMPH (TREHALOSE IS THE MAJOR CARBOHYDRATE IN THE HEMOLYMPH OF INSECTS).

-!- SIMILARITY: BELONGS TO THE AKH / HRTH / RPCH FAMILY.
            SPECIES=1.decemlineata; TISSUE=Corpora cardiaca;
MEDLINE=90160053; PubMed=2576128;
Gaede G., Kellner R.;
"The metabolic neuropeptides of the corpus cardiacum from the potato
beetle and the American cockroach are identical.";
Peptides 10:1287-1289(1989).
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01-JTB-1999 (Rel. 28, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
Hypertrehalosaemic factor II (HTF-II) (HRTH-II) (Hypertrehalosaemic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SPECIES=C.morosus; TISSUE=Corpora cardiaca; MEDLINE=87157103; PubMed=1828078; Gaede G., Rinehart K.L. Jr.; "Primary structure of the hypertrehalosaemic factor II from the corpus cardiacum of the Indian stick insect, Carausius morosus, determined by fast atom bombardment mass spectrometry."; Biol. Chem. Hoppe-Seyler 368:67-75(1987).
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Pred. No. 1.1e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       86745771A9D1A736 CRC64;
                                                                                                                                                                                                                SPECIES=B.orientalis; TISSUE=Corpora cardiaca;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       neuropeptide 11).
Carausius morosus (Indian stick insect), and
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Interpro; IPR002047; AKH.
PROSITE; PS00256; AKH: 1.
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S08996; S08996.
B44960; B44960.
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KANDINE-93129188; Pubmed=1482345;

KADLINE-93129188; Pubmed=1482345;

KADLINE-93129188; Pubmed=1482345;

Gaede G., Kellner R., Rinehart K.L. Jr., Proefke M.L.;

Gaede G., Kallner R., Rinehart K.L. Jr., Proefke M.L.;

Tar a tick fineet corpus cardiacum.";

Biochem. Biophys. Res. Commun. 189:1303-1309 (1992).

LIBOATE THE LOWIN OF TREALOSE IN THE HEMOLYMPH (TREHALOSE IS THAT BEOVATE THE LEVEL OF TREHALOSE IN THE HEMOLYMPH OF INSECTS).

LIE MAJOR CARBOHYDRATE IN THE HEMOLYMPH OF INSECTS).

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RES. SO7157; S07157.

PIR; S07157; S07157.
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01.AUG-1990 (Rel. 15, Last sequence update)
15.Unv.2002 (Rel. 41, Last annotation update)
Urease alpha subunit (EC 3.5.1.5) (Urea amidohydrolase) (Urease 63 kDa subunit) (Fragment).
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Gromphadorhina portentosa, Blattella germanica and Blatta orientalis
                     and of the stick insect Exkatosoma tiaratum assigned by tandem fast atom bombardment mass spectrometry."; Biol. Chem. Hoppe-Seyler 371:345-354 (1990).
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Hu L.-T., Nicholson E.B., Jones B.D., Lynch M.J., Mobley H.L.T.;
"Morganella morganii urease: purification, characterization, and
isolation of gene sequences.";
J. Bacteriol. 172:3073-3080(1990).
-!- CATALYTIC ACTIVITY: Urea + H(2)0 = CO(2) + 2 NH(3).
-!- COPACTOR: Binds 2 nickel ions per subunit (Potential).
-!- SUBUNIT: (ALPHA, BETA, GAMMA) (3) (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE UREASE FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
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Pred. No. 1.5e+03;
1; Mismatches 3; Indels
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25.0%; Pred. No. 2.4e+03;
cive 5; Mismatches 4;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Interpro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
Neuropeptide; Amidation; Glycoprotein.
MOD_RES 1 1 PYRROLIDO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PROSITE; PS00145; UREASE 2; PARTIAL. PROSITE; PS01120; UREASE_1; PARTIAL. Hydrolase; Metal-binding; Nickel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
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InterPro; IPR001924; UreaseA.
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50.0%;
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SEQUENCE
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VARIANT
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AKH_TABAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation - the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@lsb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ó
                                                                                                                                                                       Glycine max (Soybean).
Bukaryota, Viridiplantae; Streptophyta; Embryophyta; Tracheophyta;
Spermatophyta, Magnoliophyta; eudicotyledons; core eudicots, Rosidae;
eurosids I; Fabales; Fabaceae; Papilionoideae; Phaseoleae; Glycine.
NCBL_TaxID=3847;
                                                                                                                                                                                                                                                                                   Pearson L., Meagher R.B.;
"Diverse soybean actin transcripts contain a large intron in the 5'
untranslated leader: structural similarity to vertebrate muscle actin
                                                                                                                                                                                                                                                                                                                                                                         IN ALL EUKARYOTIC CELLS.
FUNCTION: BSSENTIAL COMPONENT OF CELL CYTOSKELETON; PLAYS AN IMPORTANT ROLE IN CYTOPLASMIC STREAMING, CELL SHAPE DETERMINATION, CELL DIVISION, ORGANELLE MOVEMENT AND EXTENSION GROWTH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                      Plant Mol. Biol. 14:513-526(1990).
-!- FUNCTION: ACTINS ARE HIGHLY CONSERVED PROTEINS THAT ARE INVOLVED IN WARIOUS TYPES OF CELL MOTILITY AND ARE UBIQUITOUSLY EXPRESSED
                                                                                                                                                                                                                                                                                                                                                                                                                          -!- SUBCELLULAR LOCATION: Cytoplasmic.
-!- MISCELLANEOUS: THERE ARE AT LEAST 16 ACTIN GENES IN SOYBEAN.
-!- SIMILARITY: BELONGS TO THE ACTIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE 17 AA; 1749 MW; 38F4970D4BEFF3C3 CRC64;
                                                                                                         01-APR-1990 (Rel. 14, Created)
01-APR-1990 (Rel. 14, Last sequence update)
15-JUL-1999 (Rel. 38, Last annotation update)
                                                                                       17 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pfam; PF00022; actin; 1.
PROSITE; PS00406; ACTINS 1; PARTIAL.
PROSITE; PS00432; ACTINS 2; PARTIAL.
PROSITE; PS01132; ACTINS ACT LIKE; PARTIAL.
Structural protein; Multigene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CPAX BOVIN STANDARD; PRT; 18 AA P22779; 01-AUG-1991 (Rel. 19, Created) 01-AUG-1991 (Rel. 19, Last sequence update)
                                                                                                                                                                                                                                                             STRAIN=cv. Wayne;
MEDLINE=91346640; Pubmed=2102831;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         EMBL; X17119; CAA34979.1; -.
PIR; S15754; S15754.
InterPro; IPR004001; Actin.
InterPro; IPR004000; Actin_like.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4; Conservative
                                                                                     STANDARD;
1 PPLSQETFSDLW 12
                        1 POISROEYGGLF 12
                                                                                                                                                 Actin 6 (Fragment).
                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
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                                                                                     SOYBN
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                                                                                    ACT6 SOY
P15986;
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                                                                        ACT6_SOYBN
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                                                           RESULT 12
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15-DEC-1998 (Rel. 37, Last annotation update)
Cytochrome P450 2A (OLF2) (Olfactive) (P52) (EC 1.14.14.1) (Fragment).
Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bosunae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE=90046758; PubMed=2813185;
Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,
Jaffe H., Raina A.K., Riley C.T., Fraser B.A., Nachman R.J.,
Vogel V.W., Zhang Y.-S., Hayes D.K.;
"Primary structure of two neuropeptide hormones with adipokinetic and
hypotrahalosemic activity isolated from the corpora cardiaca of horse
flies (Diptera).";
Proc. Natl. Acad. Sci. U.S.A. 86:8161-8164 (1989).
-I-FUNCTION: THIS HORWONE RELEASED FROM CELLS IN THE CORPORA
CARDIACA AFTER THE BEGINNING OF FLIGHT, CAUSES RELEASE OF
DIGTYCERNDES FROM THE FAT BODY AND THEN STIMULATES THE FLIGHT
MUSCLES TO USE THESE DIGLYCERIDES AS AN ENERGY SOURCE.
-I-SIMILARITY: BELONGS TO THE AKH / HRTH / RPCH FAMILY.
PIR; A33995; A33995.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bukaryota; Metazoa; Arthopoda; Mandibulata; Pancrustacea; Hexapoda;
Insecta; Pterygota; Neoptera; Endopterygota; Diptera; Brachycera;
Tabanomorpha; Tabanidae; Tabanus.
NCBI_TaxID=7207;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PIR; A35704; A35704.
InterPro; IPR001128; Cytochrome P450.
PROSITE; P500086; CYTOCHROME_P450; PARTIAL.
Oxidoreductase; Monocoxygenase; Blectron transport; Membrane; Heme;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -!- CATALYTIC ACTIVITY: RH + reduced flavoprotein + O(2) = ROH + oxidized flavoprotein + H(2)O.
-!- SUBCELLULAR LOCATION: Membrane-bound. Endoplasmic reticulum.
-!- SIMILARITY: BELONGS TO THE CYTOCHROME P450 FAMILY.
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01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
Adipokinetic hormone (AKH) (Dipteran corpora cardiaca factor I)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20.0%; Score 21; DB 1; Length 18; 36.4%; Pred. No. 2.9e+03; ive 2; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18 AA; 2058 MW; F80746F76CCD77FF CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ACIDS, AND XENOBIOTICS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Tabanus atratus (Horse fly).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 36.4%,
A; Conservative
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SPECIES=P.americana;
MEDLINE=84298179; PubMed=6591205;
Scarborough R.M., Jamieson G.C., Kalish F., Kramer S.J., McEnroe G.A.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Primary structures of the cockroaches Leucophaea maderae, the corpora cardiaca of the cockroaches Leucophaea maderae, dromphadorhina portentosa, Blattella germanica and Blatta orientalis and of the stick insect Extatosoma tiaratum assigned by tandem fast actom bombardment mass spectrometry.", Biol. Chem. Hoppe-Seyler 371:345-354 (1990).

-:- FUNCTION: HYPERTREHALOSAEMIC FACTORS ARE NEUROPEPTIDES THAT ELEVATE THE LEVEL OF TREHALOSE IN THE HEMOLYMPH (TREHALOSE IS 'THE MAJOR CARBOHYDRATE IN THE HEMOLYMPH OF INSECTS).
                                                                                                                                                                                                                                                                                                                                                                                                   01-FEB-1994 (Rel. 28, Last sequence update)
01-FEB-1994 (Rel. 28, Last annotation update)
Hypertrehalosaemic factor I (Neuropeptide M.)
(PeA-CAH-I) (LeD-CC-I) (Hypertrehalosaemic neuropeptide I).
Periplaneta americana (American cockroach),
Leptinotaras decemilineata (Colorado potato beetle), and
Blatta orientalis (Oriental cockroach).
Eukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
Insecta; Perrygota; Neoptera; Orthopteroidea; Dictyoptera; Blattaria;
Blattoidea; Blattidae; Periplaneta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SPECIES=L.decemlineata; TISSUE=Corpora cardiaca;
MEDLINE=90160053; PubMed=2576128;
Gaede G., Kellner R.;
"The metabolic neuropeptides of the corpus cardiacum from the potato
beelle and the American cockroach are identical.";
Peptides 10:1287-1289(1989).
                                                                                                                                                                         0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Structures of two cockroach neuropeptides assigned by fast atom bombardment mass spectrometry."; Biochem. Biophys. Res. Commun. 124:350-358(1984).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SPECIES=P.americana;
MEDLINE=85046530; PubMed=6548628;
Witten J.L., Schaffer M.H., O'Shea M., Cook J.C., Hemling M.E.,
Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Miller C.A., Schooley D.A.;
"Isolation and primary structure of two peptides with
cardioacceleratory and hyperglycemic activity from the corpora
cardiaca of Periplaneta americana.";
Proc. Natl. Acad. Sci. U.S.A. 81:5575-5579 (1984).
                      Neuropeptide; Amidation; Flight.

MOD RES 1 1 PYRROLIDONE CARBOXYLIC ACID.

MOD_RES 8 AMIDATION
                                                                                                                                Score 20; DB 1; Length 8; Pred. No. 1.1e+05; Mismatches 3; Indels
                                                                          MOD_RES 8 8 AMIDATION.
SEQUENCE 8 AA; 949 MW; 86786771A9D1A736 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SPECIES=B.orientalis; TISSUE=Corpora cardiaca;
                                                                                                                                                                                                                                                                                                                                                   8 AA.
                                                                                                         Query Match 19.0%; Score 20; UB A Best Local Similarity 50.0%; Pred. No. 1.1e+
                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=90253659; PubMed=2340112;
Gaede G., Rinehart K.L. Jr.;
                                                                                                                                                                                                                                                                                                                                                                                         13-AUG-1987 (Rel. 05, Created)
                                                                                                                                                                                                                                                                                                                                                     STANDARD;
InterPro; IPR002047; AKH.
PROSITE; PS00256; AKH; 1.
                                                                                                                                                                                                                 5 QETFSDLW 12
                                                                                                                                                                                                                                                      1 QLTFTPGW 8
                                                                                                                                                                                                                                                                                                                                                     PERAM
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CC -1- SIMILARITY: BELONGS TO THE AKH / HRTH / RPCH FAMILY.

DR PIR; A05169; A05169.

DR PIR; A49862; A49823.

DR PIR; A449862, A44986.

DR INCEPPO: IPROACO47; AKH.

DR PKOSITE; PS00256, A4496.

KW Neuropeptide; Amidation.

FT MOD_RES 8 8 8 AMIDATION.

FT MOD_RES 991 MW; 86745775B9C452D6 CRC64;

Query Match 19.0%; Score 20; DB 1; Length 8;

Best Local Similarity 50.0%; Pred. No. 1.1e+05;

Matches 4; Conservative 0; Mismatches 4; Indels 0; Gaps

Qy 5 GETFSDLW 12

Db 1 QVNFSDWW 8

Search completed: February 12, 2003, 10:44:24

Job time: 30 secs
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

February 12, 2003, 10:42:53; Search time 46 Seconds

(without alignments) 39.708 Million cell updates/sec

US-09-403-440A-2 Perfect score:

1 PPLSQETFSDLWKLLPENG 19 BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table: Sequence:

283224 seqs, 96134422 residues Searched:

3433 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 19

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR_73:* Database :

pirl:* pir2:* pir3:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description		polyprotei	1 (hypotherical proce	himograph elonga	dhannaria 18 1	adinoting to homo	adiportieric normo	0	procein Kinase (EC	ಹ	prot	, H	7		lymnaDFamide 5 - g	L-2,4-diaminobutyr	insulin-like growt	hydroxyproline-ric	Ig H chain V-D-J r	T cell receptor al	shikimate 5-dehydr	hypertrehalosemic	adipokinetic hormo	neuronentide Led-	hymertrebalogomia	hmontropy	TAPET CIETOSEMIC	r)	214A excantigen (v beta-D-galactosida
SUMMARIES	0	B61409	A59396	A35105	PA0110	A31571	S41747	S11545	A24244	520410	570615	516324	532471	4444	27 77 70	4/4700	0117	544854	0.0517	S59481	PH1625	PH1788	140062	8996	B49823	44960	JC1416	09138	53789	309	343
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de	Query Match	32.4	24.8	23.8	23.8	22.9	22.9	21.9	21.9	21.9	21.0	21.0	21.0	21.0	21.0	21.0	0.10			2 1 0	0.0	0.0				70.0	20.0	20.0	20.0	20.0	20.0
	Score	34	26	25	25	24	24	23	23	23	22	22	22	22	22	22	22	22	2.0	21.5	11.0	21:5		1 6	1 .	77	21	21	21	21	21
	Result No.	-	7	m	4	Ŋ	Q	7	80	o	10	11	12	13	14	15	16	17	8	9 6	6	2 5	22	3 6	9 6	7 L	52	56	27	28	59

histone Hi.c - mou histone Hia - mous	ulpeptidyi-peptida urease (EC 3.5.1.5 7 alpha-hydroxy-4-	superoxide dismuta actin 6 - soybean	cytochrome P450 ol TCR delta chain V-	Cysteine proteinas Ig H Chain V-D-J r hynothetical arato	hypertrehalosemic	neuropeptide Led-C adipokinetic hormo
13 2 B35245 13 2 A35245 13 2 B56864	15 2 A35389 15 2 A45103	16 2 E58503 17 2 S15754 16 3 %37535	18 2 A35/04 19 2 I49037 13 2 PC4391	14 2 PH1627 18 2 I52623	8 2 S08995 8 2 A49823	8 2 A44960 8 2 A33995
21 20.0 21 20.0 21 20.0	21 20.0	21 20.0			20 19.0 20 19.0	20 19.0 20 19.0
30 31 32	333 143	36	388	40	4 4 2 2 2	4. 4. 4. 7.

ALIGNMENTS

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Genome polyprotein (clone L3/S2) - Skalica virus (fragment)
C;Species: Skalica virus
C;Species: 19-Mar-1997 #sequence_revision 19-Dec-1997 #text_change 17-Nov-2000
C;Accession: B61409
R;Guirakhoo, F.; Heinz, F.X.; Mandl, C.W.; Holzmann, H.; Kunz, C.; Gresikova, M.
J. Gen. Virol. 72, 333-338, 1991
A;Title: The relationship between the flaviviruses Skalica and Langat as revealed by most A;Reference number. A61409; MUID:91132129; PMID:1847173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
                                                                                                                                                                                                                                                                                                                                        A)Status: preliminary; not compared with conceptual translation A;Molecule type: genomic RNA A;Residues: 1-19 <GUI> C;Superfamily: yellow fever virus genome polyprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match

32.4%; Score 34; DB 2; Length 19;
Best Local Similarity 41.7%; Pred. No. 81;
Matches 5; Conservative 3; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 PPLSQETFSDLW 12
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A59396 Tha p 1 - Thaumetopoea pityocampa (fragment) Cispeciaes: Thaumetopoea pityocampa Cispeciaes: Thaumetopoea pityocampa Cibete: 03-Jun-2002 #sequence_revision 03-Jun-2002 #text_change 03-Jun-2002 CiAccession: A59396 R:Moneo, I. A:Description: Isolation and characterization of a major allergen of the pine procession A:Reference number: A59396 2 PPVTYRTGTDCW 13 RESULT 2 g

ö 0; Gaps 24.8%; Score 26; DB 2; Length 18; 38.5%; Pred. No. 1.28+03; tive 3; Mismatches 5; Indels A; Note: IgE-binding protein, major allergen 5; Conservative A; Experimental source: L5 A;Status: preliminary A;Molecule type: protein A;Residues: 1-18 <MON> Local Similarity Query Match Best Loca Matches

6 ETFSDLWKLLPEN 18 2 ETYSDKYDTIDVN 14 a ò

RESULT 3

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dipokinetic hormone - bollworm

N.Alternate names: Hez-AKH

N.Alternate names: Hez-AKH

C.Species: Heliothis zea (bollworm, corn earworm, tomato fruitworm)

C.Species: Heliothis zea (bollworm, corn earworm, tomato fruitworm)

C.Accession: A24244

R.Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Holman, G.M.; Wagner, R.M.; Ridgwa Blochem. Blophys. Res. Commun. 135, 622-628, 1986

A.Jitle: Isolation and primary structure of a peptide from the corpora cardiaca of Heli
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Accession: A24244
A;Molecule type: protein
A;Residues: 1-9 < AAF.
C;Superfamily: adipokinetic hormone
C;Superfamily: adipokinetic hormone
C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; Fyroglutami
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Superfamily: adipokinetic hormone C;Superfamily: adipokinetic hormone neuropeptide; pyroglutamil C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamil F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental P;8/Modified site: amidated carboxyl end (Trp) #status experimental
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Accession: S11545
R;Gaede, G.; Wilps, H.; Kellner, R.
Biochem. J. 269, 309-313, 1990
A;Title: Isolation and structure of a novel charged member of the red-pigment-concentral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               adipokinetic hormone - nestling-sucking blowfly
C;Species: Protophormia terraenovae (nestling-sucking blowfly)
C;Date: 17-Apr-1993 #sequence_revision 17-Apr-1993 #text_change 31-Oct-1997
                                                                                                                                                                                                            C;Species: Solanum tüberösum (potatö)
C;Date: 19-Mar-1997 #sequence_revision 29-Aug-1997 #text_change 07-May-1999
                                                                                                                                                                                                                                                                                                      R;Burt, W.J.E.; Leaver, C.J.
FEBS Lett. 339, 139-141, 1994
A;Title: Identification of a chaperonin-10 homologue in plant mitochondria.
A;Reference number: S41747; MUID:94148071; PMID:7906228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21.9%; Score 23; DB 2; Length 8; 62.5%; Pred. No. 2.8e+05; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22.9%; Score 24; DB 2; Length 11; 83.3%; Pred. No. 1.5e+03; ative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Reference number: S11545; MUID: 90351345; PMID: 2386478
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A, Experimental source: mitochondrion C, Keywords: mitochondrion; molecular chaperone
                                                                                                                                                                                           chaperonin 10 homolog - potato (fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Molecule type: protein A;Residues: 1-8 <GAE>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
les 5; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: protein
A; Residues: 1-11 < BUR>
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 QLTFSPDW 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  erraenovae (Diptera)
| ||| |
1 QLTFSSGW 8
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                                                                                                                                                                                                                                                                                            C;Accession: S41747
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14 LLPENG 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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C;Species: Heliothis zea (bollworm, corn earworm, tomato fruitworm)
C;Species: Heliothis zea (bollworm, corn earworm, tomato fruitworm)
C;Species: Jo-Unn-1989 #sequence_revision 23-Mar-1995 #text_change 31-Oct-1997
C;Accession: A31571
R;Jaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Bird, T.G.; Tseng, C.M.; Zhang, Y.S
B;Jaffe, H.; Raina, Res. Commun. 155, 344-350, 1988
A;Title: Isolation and primary structure of a neuropeptide hormone from Heliothis zea will reference number: A31571; MUID:88326324; PMID:3415690
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C.Superization of adipokinetic hormone C.Keywords: anidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamid C;Keywords: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamid F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F;10/Modified site: amidated carboxyl end (Asn) #status experimental
                           hypothetical protein - Neurospora crassa mitochondrion (fragment)
hypothes: mitochondrion Neurospora crassa
(;Species: mitochondrion Neurospora crassa
C;Date: 07-Sep-1990 #sequence_revision 07-Sep-1990 #text_change 07-Dec-1999
C;Accession: A35105
E;Accession: A35105
A;Title: A site-specific self-cleavage reaction performed by a novel RNA in neurospora fa;Reference number: A35105; MUID:90263093; PMID:2160856
A;Accession: A35105
A;Eterus: preliminary; not compared with conceptual translation
A;Molecule type: DNA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         translation elongation factor eBF-1 beta' chain - Arabidopsis thaliana (fragment) c;Species: Arabidopsis thaliana (mouse-ear cress) c;Date: 07-Apr-1995 #sequence_revision 26-May-1995 #text_change 26-Apr-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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22.9%; Score 24; DB 2; Length 10;
Best Local Similarity 62.5%; Pred. No. 1.38+03;
Matches 5; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           23.8%; Score 25; DB 2; Length 15; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23.8%; Score 25; DB 2; Length 14; 54.5%; Pred. No. 1.3e+03; tive 0; Mismatches 5; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CiAccession: PA0110
Rikamo, M.; Kawakami, T.; Tsugita, A.
submitted to JIPID, March 1995
A;Reference number: PA0109
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Matches 5; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C; Keywords: mitochondrion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: protein A; Residues: 1-15 < KAM>
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Query Match

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hypothetical protein 2 - Arabidopsis thaliana (Species: Arabidopsis thaliana (Mouse-ear cress)
C;Species: Arabidopsis thaliana (Mouse-ear cress)
C;Date: 2.1-Nov-1993 #sequence_revision 12-May-1995 #text_change 21-Jul-2000
C;Accession: S16324
R;Ruberti, I.; Sessa, G.; Lucchetti, S.; Morelli, G.
R;Ruberti, I.; Sessa, G.; Lucchetti, S.; Morelli, G.
A;Title: A novel class of plant proteins containing a homeodomain with a closely linked A;Reference number: S16323; MUID:91266907; PMID:1675603
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cispecies: Lymnaea stagnals (great pond snail)
Cibate: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
Cipate: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
Cipatesion: 832472
RiJohnsen, A.H.; 816-879, 1993
Aitile: LymnaDFamides, a new family of neuropeptides from the pond snail, Lymnaea stagnareference number: 832471; MUID:93238777; PMID:8477756
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C.Species: Lymnaea stagnalis (great pond snail)
C.Date: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
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A,Residues: 1-13 «JOH»
A,Cross-references: PIDN:AAB26362.1; PID:g299829
A,Experimental source: ganglia
C,Keywords: amidated carboxyl end, neuropeptide
F,13/Modified site: amidated carboxyl end (Phe) #status predicted
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A/Residues: 1-13 <JOH>
A/Residues: 1-13 <JOH>
A/Cross-references: PIDN:AAB26363.1; PID:g299830
A/Experimental source: ganglia
C;Keywords: amidated carboxyl end; neuropeptide
F;13/Modified site: amidated carboxyl end (Phe) #status predicted
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     21.0%; Score 22; DB 2; Length 13; 50.0%; Pred. No. 3.5e+03; Live 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21.0%; Score 22; DB 2; Length 8; 80.0%; Pred. No. 2.8e+05; Live 1; Mismatches 0; Indels
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Matches 4; Conserva
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                             A; Residues: 1-8 < RUB>
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3 YKLLP 7
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PEBS Lett. 298, 33-35, 1992
A;Title: Phosphorylation of cytochrome b6 by the LHC II kinase associated with the cytochrone a;Reference number: S20410; MUID:92183823; PMID:1544419
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Grades Streptomyces sp. (Chainia sp. NCL 82.5.1) (fragmen N.Alternate names: xylanase (S. 3.2.1.8) - Streptomyces sp. (Chainia sp. NCL 82.5.1) (fragmen N.Alternate names: xylanase Streptomyces sp. A.Variety: Chainia sp. NCL 82.5.1 (fragmen A.Variety: Chainia sp. NCL 82.5.1 (fragmen C.) Date: 19-Mar-1998 #sequence_revision 17-Apr-1998 #text_change 07-May-1999 R.Rao. M.; Khadilkar, S.; Bandivadekar, K.R.; Deshpande, V. Biochem. J. 316, 771-775, 1996 A.Title: Structural environment of an essential cysteine residue of xylanase from Chainia A.Reference number: S70615; MUID: 96265041; PMID: 8670151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Description: is responsible for the regulation of energy distribution between photosys A; Note: does not exhibit redox-controlled activation C; Keywords: chloroplast; light-harvesting complex; membrane-associated complex; phosphoc
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                                                                                                                                                                                                                                                                                                                                                                                        Drotein kinase (BC 2.7.1.37) - spinach chloroplast (fragment)
N;Alternate names: LHCII protein kinase
C;Species: chloroplast Spinacia oleracea (spinach)
C;Date: 19-Mar-1997 #sequence_revision 11-Jun-1999 #text_change 11-Jun-1999
C;Accession: S20410
                                                                                                                                                            Gaps
F:1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental F:9/Modified site: amidated carboxyl end (Gly) #status experimental
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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40.0%; Pred. No. 2.9e+03;
live 2; Mismatches 4; Indels
                                                                                        21.9%; Score 23; DB 2; Length 9; 50.0%; Pred. No. 2.8e+05; Live 1; Mismatches 3; Indels
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A/Molecule type: protein
A/Residues: 1-5 <RAO>
A/Experimental source: Chainia sp. strain NCL 82.5.1
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                                                        Query Match
Best Local Similarity 50.00
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Matches 4; Conserv
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C; Function:
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Cispecies: Lymnaea stagnalis (great pond snail)
Cjacte: 19-Mar-1997 #sequence_revision 19-Mar-1997 #text_change 20-Aug-1999
CjAccession: 832475
RjOhnsen, A.H.; Rehfeld, J.F.
RjOhnsen, A.H.; Rehfeld, J.P.
A.Fitle: LymnaPanides, a new family of neuropeptides from the pond snail, Lymnaea stagn A; Reference number: 832471; MUD:93238777; PMID:8477756
                                                                                                                                                                                                                                                                                                                                                                          tymmatronius = 1 Strain Johnson 2018 (great pond snail)
C;Species: Lymnaea stagnalis (great pond snail)
C;Species: Lymnaea stagnalis (great pond snail)
C;Species: Lymnaea stagnalis (great pond snail)
C;Accession: S32474
R;Johnsen, A.H.; Rehfeld, J.F.
Bur. J. Biochem. 213, 875-879, 1993
A;Title: LymnabFanides, a new family of neuropeptides from the pond snail, Lymnaea stagn A;Reference number: S32474
A;Accession: S32474
A;A
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A;Experimental source: ganglia
C;Keywords: amidated carboxyl end; neuropeptide
F;13/Modified site: amidated carboxyl end (Phe) #status predicted
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21.0%; Score 22; DB 2; Length 13;
Best Local Similarity 50.0%; Pred. No. 3.5e+03;
Matches 4; Conservative 1; Mismatches 3; Indels
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1: /cgn2_6/ptodata/1/iaa/5A_COMB.pep:*
2: /cgn2_6/ptodata/1/iaa/5B_COMB.pep:*
3: /cgn2_6/ptodata/1/iaa/6A_COMB.pep:*
4: /cgn2_6/ptodata/1/iaa/6B_COMB.pep:*
5: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
6: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-081-975-10

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US-08-245-50A-1

US-08-390-546-1

US-08-390-546-1

US-08-390-546-1

US-08-390-515A-1

US-08-390-515A-1

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US-08-390-515A-1

US-08-390-515A-1

US-08-390-515A-1

US-08-697-221-19

US-08-697-221-120

US-08-697-221-221

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                                                                                                                                                                                                                                                                                                                                              262574 seqs, 29422922 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                        OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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seq length: 200000000
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Match Length DB
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Maximum DB
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APPLICANT: Acelin, William
APPLICANT: Jost, Christine
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODIS
TITLE OF INVENTION: ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES:
ADDRESSEE: Nixon Peabody LLP
STEEFT: 101 Federal Street
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COMPUREY. READABLE FORM:
MEDLIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: JESSE OF WINGOWS
SOFTWARE: FASTSEO for WINGOWS
SOFTWARE: FASTSEO for WINGOWS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATYOMEY/AGENT INFORMATION:
NAME: EISENStein, Rohald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
          US-08-130-515C-7

US-08-130-515C-8

US-08-130-515C-9

US-08-131-357-2

US-08-130-515A-6

US-08-130-515A-7

US-08-130-515A-9

US-08-130-5111

US-08-130-51114

US-08-130-51114

US-08-130-51114

US-08-130-51114
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Patent No. 6451979
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LENGTH: 20 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 617-345-1300
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US-09-081-975-8
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ADDRL.
STREET: LV.
TTV: Boston
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US-09-081-975-8
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STREET:
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GENERAL INFORMATION:
APPLICANT: Realin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE, OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE, OF INVENTION: ANTIBODIES
TITLE OF SEQUENCES: 27
                                                                                                                                                                                                                          APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
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94.3%; Score 99; DB 4; Length 20;
Best Local Similarity 100.0%; Pred. No. 1.6e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IEM Compatible
COMPUTER: IEM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastERG for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTOMNEY/AGENT INFORMATION:
NAME: Elsenstein, Ronald I
REGISTRATION NUMBER: 30,628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-081-975-11
; Sequence 11, Application US/09081975
; Patent No. 6451979
                                                                                                                                                    Sequence 10, Application US/09081975
Patent No. 6451979
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         47400
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STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 474
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 10:
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       1 PPLSOETFSDLWKLLPEN 18
                            2 PPLSQETFSDLWKLLPEN 19
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: linear
US-09-081-975-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     USA
                                                                                                                                                                                                                                                                                                                                                                                                                                     Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        02110
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                                                                                                                                        US-09-081-975-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 3
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APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match

Best Local Similarity 100.0%; Pred. No. 1.6e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: Fast-SEQ for Windows
SOFTWARE: TAST-SEQ for Windows
SOFTWARE: 12-MA-1998
CLASSIPICATION UNDER: US/09/081,975
FILING DATE: 12-MA-1998
CLASSIPICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                   COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COPERATING SYSTEM: Windows
SOFTWARE: FESTESEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION BATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 sequence 16, Application US/09081975; Patent No. 6451979; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                      NAME: EISENStein, Rohald I
REGISTRATION NUMBER: 30,628
REFERIOR/DOCKET NUMBER: 4740(
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nixon Peabody LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 101 Federal Street
CITY: Boston
101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 PPLSOETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                       Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; STRANDEDNESS: single
; TOPOLOGY: linear
US-09-081-975-11
                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acid
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                        Boston
                                                                   USA
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Gaps

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Query Match 94.3%; Score 99; DB 1; Length 64; Best Local Similarity 100.0%; Pred. No. 5.9e-09; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: BURRELL, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: VGELSTBIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
ADDRESSEE: BANNER, BIRCH, MCKIE AND BECKETT
STREET: 1001 G STREET, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94.3%; Score 99; DB 1; Length 64; 100.0%; Pred. No. 5.9e-09; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READBLE FORM:
MODIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,546
FILING DATE: O'-APR 1993
CLASSIFICATION NUMBER: 33,141
REFERENCE/DOCKET NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.42798
TELEPHONE: 202-508-9100
TELEPHONE: 202-508-929
                                                                                                                                                                                                                                                                                                            RESULT 6
US-08-390-546-1
S-08-390-546-1
S equence 1, Application US/08390546
Patent No. 5606044
GENERAL INFORMATION:
                                                                                                                                                                                         1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                    12 PPLSQETFSDLWKLLPEN 29
                    CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         N-terminal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WASHINGTON
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FRAGMENT TYPE: NORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: WASHI
STATE: D.C.
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US-08-245-500A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-390-546-1
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APPLICANT: BURRELL, MARILEE
APPLICANT: HILL, DAVID B.
APPLICANT: KINZLER, KENDETH W.
APPLICANT: KINZLER, KENDETH W.
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN TITLE OF SEQUENCES: 5
CORRESPONDENCES: 5
CORRESPONDENCES: 5
ADDRESSE: BANNER, BIRCH, MCKIE AND BECKETT
STREET: 1001 GTREET, N.W.
                                                                                                                                                                                                                                                                                                                                                                            Query Match 94.3%; Score 99; DB 4; Length 20; Best Local Similarity 100.0%; Pred. No. 1.6e-09; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 2001

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC COMPATIBLE

COMPATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/245,500A

FILING DATE: 07-APR-1993

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:

NAME: KAGAN, SARAH A.

REGISTRATION NUMBER: 32,141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.42798
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1, Application US/08245500A
Patent No. 5550023
NAWE: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 4740
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                         information for SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
ENGTH: 20 amino acids
TYPE: amino acids
STRANDENESS: single
TOPOLOGY: linear
US-09-081-975-16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPAX: 202-508-9299
TELEX: 197430 BBMB UT
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 PPLSOETFSDLWKLLPEN 18
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ORIGINAL SOURCE:
ORGANISM: Homo sapiens
POSITION IN GENOME:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 64 amino acids
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STRANDEDNESS: single
TOPOLGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-245-500A-1
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ZIP: 20001
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                     ZIP: 20001
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FRAGMENT TYPE: N-terminal ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein
CORRESPONDENCE ADDRESS: ADDRESSE: BANNER, B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     linear
                                                               WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: amino a
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-557-393-1
                                                                                                          COUNTRY:
                                                                                     STATE:
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94.3%; Score 99; DB 1; Length 64;
Best Local Similarity 100.0%; Pred. No. 5.9e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1, Application US/08557393
Patent No. 5702903
GENERAL INFORMATION:
APPLICANT: HILL, DAVID E.
APPLICANT: HILL, DAVID E.
APPLICANT: HILL, DAVID E.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF 'SEQUENCES: 5
                                                                                                     APPLICANT: BURRELL, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: HILL, DAVID E.
APPLICANT: KINELER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN
WIMBER OF INVENTION: HUMAN TUMORS
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/390,479A
FILING DATE: US/08/390,479A
FILING DATE: US/08/390,479A
FILING DATE: US/08/390,479A
ATYONEN'AGENT INFORMATION:

NAME: REARRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.48992
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SOURCE: US/08/999
TELESTA: 197430 BBMB UT
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                ADDRESSEE: BANNER & WITCOFF, LTD. STREET: 1001 G STREET, N.W. CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                          Sequence 1, Application US/08390479A Patent No. 5618921 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 PPLSOETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 64 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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                                                                                                                                                                                                                                                                                                                              STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                           20001
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US-08-557-393-1
         RESULT 7
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ADDRESSES, BANKER, BIRCH, MCKIE AND BECKETT
STREET: 100.1 G STREET, N.W.
CITY: WASHINGTON
CITY: WASHINGTON
COUNTRY: D.C.
COUNTRY: USA
C
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APPLICANT: BURELL, MARILEE
APPLICANT: BURELL, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF SEQUENCES: 9
CORRESPONDENCE ADDRESS:
ADDRESSEE: BANNER, BIRCH, MCKIE AND BECKETT
STREET: 1001 G STREET, N.W.
CITY: WASHINGTON
STRATE: D.C.
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 94.3%; Score 99; DB 1; Length 64; Best Local Similarity 100.0%; Pred. No. 5.9e-09; Matches 18; Conservative 0; Mismatches 0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOGTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,515A
FILING DATE: 07-APR-1993
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 01107.42798
TELEPHONE: 202-508-9100
TELEPHONE: 202-508-9100
TELEPRAX: 202-508-910
TELEPAX: 202-508-929
TELEX: 197430 BBMB UT
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 64 amino acids
TYPE: amino acid
STRANDEDDRES: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 1, Application US/08390515A; Patent No. 5756455
              TELEPHONE: 202-508-9100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens

POSITION IN GENOME:

CHROMOSOME/SEGMENT: 17q

US-08-390-517A-1
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                                                                                                                                                                                                                                                                                                                                                                                                                            N-terminal
                                                                                                                                                                               LENGTH: 64 amino acids
                                       TELEFAX: 202-508-9299
TELEX: 197430 BBMB UT
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                       single
                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
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MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                    linear
                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: si:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
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D.C.
USA
                                                                                                                                                                                                                                                                                                                                                                                                                         FRAGMENT TYPE: NORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ANTI-SENSE: NO
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US-08-390-515A-1
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APPLICANT: BURRELL, MARILEE

APPLICANT: HILL, DAVID E.

APPLICANT: HILL, DAVID E.

APPLICANT: WOGELSTEIN, BERT

TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN

TITLE OF INVENTION: AMPLIFICATION OF HUMAN MDM2 GENE IN

TITLE OF INVENTION: BIRRT

TITLE OF INVENTION: WAPLIFICATION OF HUMAN MDM2 GENE IN

TITLE OF INVENTION: BIRRT

TITLE OF INVENTION: WAPLIFICATION OF HUMAN MDM2 GENE IN

TITLE OF INVENTION: WAPLIFICATION OF HUMAN MDM2 GENE IN

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TITLE OF INVENTION: WAPLIFICATION OF HUMAN MDM2 GENE

TITLE OF INVENTION: WAPLIFICANT OF WAPLIFICATION OF HUMAN MDM2 GENE

TITLE OF INVENTION: WAPLIFICATION OF HUMAN MDM2 GENE

TITLE OF INVENTION OF HUMAN MDM2 GENE

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TITLE OF INVENTION OF HUMAN MDM2 GENE

TITLE OF IN
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                                              OPERATION SYSTEM: TC-DUS/MS-LUS
SOFTWARE: PATCHIEN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/390,516C
FILING DATE: 07-APR-1993
CLASSIFICATION: 530
ATTONNEY/AGENT INPORMATION:
NAME: REGISTRATION NUMBER: 32,141
REFERENCE/POCKET NUMBER: 32,141
REFERENCE/POCKET NUMBER: 32,141
REFERENCE/POCKET NUMBER: 32,141
RELEFAX: 202-508-9100
TELLEFAX: 202-508-929
TELLEFAX: 202-508-929
TELLER: 197430 BBMB UT
SEQUENCE CHARACTER/STICS:
SEQUENCE CHARACTER/STICS:
TURDAMATION FOR SEQ ID NO: 1:
SEQUENCE CAPACTER/STICS:
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
SOFTWARE: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
APPLICATION NUMBER: US/08/390,517A
FILING DATE: 07-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/POCKET NUMBER: 32,141
REFERENCE/POCKET NUMBER: 01107.42798
TELECOMMUNICATION INFORMATION:
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ANTI-SENSE: NO
FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Homo sapiens
POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 64 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
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Matches 18; Conservative
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US-09-170-159A-1
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                                                                                                                                                                                                                                                                Query Match

94.3%; Score 99; DB 1; Length 64;
Best Local Similarity 100.0%; Pred. No. 5.9e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: BURRELL, MARILEE
APPLICANT: HILL, DAVID E.
APPLICANT: KINZLER, KENNETH W.
APPLICANT: VOGELSTEIN, BERT
TITLE OF INVENTION: HUMAN TUMORS
NUMBER OF SEQUENCES: 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/801,718
FILING DATE: 14-FEB-1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/390,515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ...ukeSSEE: BANNER, BIRCH, MCKIE AND BECKETT STREET: 1001 G STREET, N.W. CITY: WASHINGTON STATE: D.C. TATE: D.
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FILING DATE: 07-APR-1993
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 32,141
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-508-9100
TELEFAX: 202-508-910
TELEFX: 197430 BBMB UT
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/08801718
Patent No. 5858976
GENERAL INFORMATION:
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100.0%;
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CHROMOSOME/SEGMENT: 17q
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                                                                             ORGANISM: Homo sapiens
POSITION IN GENOME:
CHROMOSOME/SEGMENT: 17q
                                                                                                                                                                                                                                                                                                                                                                                                                               1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12 PPLSQETFSDLWKLLPEN 29
            N-terminal
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STRANDEDNESS: Si
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Best Local Similarity
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D.C.
USA
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        FRAGMENT TYPE: NORIGINAL SOURCE:
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HILL, DAVID B.
KINZLER, KENNETH W.
VOGELSTEIN, BERT
VOGELSTEIN, BERT
HUMAN TUMORS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 94.3%; Score 99; DB 4; Length 64; Best Local Similarity 100.0%; Pred. No. 5.9e-09; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: BANNER, BIRCH, MCKIE AND BECKETT STREET: 1001 G STREET, N.W.
CITY: WASHINGTON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/170,159A
FILING DATE: 13-Oct-1998
CLASSIFICATION: «Unknown»
ATTORNEY/AGENT INFORMATION:
NAME: KAGAN, SARAH A.
REGISTRATION NUMBER: 32,141
REFERENCE/DOCKET NUMBER: 01107.42798
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CHROMOSOME/SEGMENT: 17q ; SEQUENCE DESCRIPTION: SEQ 1D NO: 1: US-09-170-159A-1
                                                                                                                                                     Sequence 1, Application US/09170159A; Patent No. 6399755; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 3, Application US/09414436
; Patent No. 6294384
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: 202-508-9100
TELEFAX: 202-508-9299
TELEX: 197430 BBMB UT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
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FRAGMENT TYPE: N-terminal
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: protein
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 PPLSQETFSDLWKLLPEN 18
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                                   12 PPLSQETFSDLWKLLPEN 29
1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: D.C.
COUNTRY: USA
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APPLICANT: Dell'Acqua, Giorgio
APPLICANT: Dell'Acqua, Giorgio
APPLICANT: Mann, Michael J.
APPLICANT: Mann, Michael J.
APPLICANT: Dzau, Victor J.
TITLE OF INVENTION: Compositions and Methods Based Upon an Isoform of p53
FILE REFERENCE: p53f
CURRENT APPLICATION NUMBER: US/09/414,436
CURRENT PILING DATE: 1999-10-07
EARLIER FILING DATE: 1999-10-07
SARLIER FILING DATE: 1999-10-09
SOFTWARE: PALENTIN NUMBER: 60/103,849
EARLIER FILING DATE: 1999-10-09
SOFTWARE: PALENTIN VE: 2.0
SEQ ID NO 3
LENGTH: 261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          % Sequence 17, Application US/08697221

Sequence 17, Application US/08697221

Fatent No. 584708:

Patent No. 584708:

APPLICANT: Halazonetis, Thanos D.

TITLE OF INVENTION: Modified p53 Constructs and Uses

TITLE OF INVENTION: Therefor

NUMBER OF SEQUENCES: 33

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Howson and Howson
STREET: Spring House Corporate Cntr., PO Box 457
CITY: Spring House
STATE: Pennsylvania
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 19477
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/697,221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/004,802
FILING DATE: 22-SEP-1995
ATTORNEY AGENT INFORMATION:
NAME: Kodroff, Cathy A.
REGISTRATION NUMBER: 33,980
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       WST64AUSA
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TELECOMMUNICATION INFORMATION:
TELEFRAK: 215-540-9206
TELEFAX: 215-540-5818
INFORMATION FOR SEQ ID NO: 17:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 363 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 PPLSQETFSDLWKLLPEN 18
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MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                      TYPE: PRT
CORGANISM: Homo sapiens
US-09-414-436-3
GENERAL INFORMATION:
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1 PPLSQETFSDLWKLLPEN 18
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Sequence 25, Appl
Sequence 26, Appl
Sequence 27, Appl
Sequence 28, Appl
Sequence 38, Appl
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Sequence 21, App
Sequence 27, Appl
Sequence 27, Appl
Sequence 286, App
Sequence 250, App
Sequence 5, App
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Sequence 3, Appli
Sequence 9, Appli
Sequence 4, Appli
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Sequence 56463, A
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Sequence 118, App
Sequence 9, Appli
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Sequence 220, App
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Sequence 3137, Ap
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                                                                                   February 12, 2003, 10:35:07; Search time 19 Seconds (without alignments) 83.787 Million cell updates/sec
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Sequence 28,
Sequence 307,
Sequence 276
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1: /cgn2 6/ptodata/1/paa/USO6_NEW_COMB.pep:*

2: /cgn2 6/ptodata/1/paa/USO6_NEW_COMB.pep:*

3: /cgn2 6/ptodata/1/paa/USO7_NEW_COMB.pep:*

4: /cgn2 6/ptodata/1/paa/USO9_NEW_COMB.pep:*

5: /cgn2 6/ptodata/1/paa/USO9_NEW_COMB.pep:*

5: /cgn2 6/ptodata/1/paa/USO3_NEW_COMB.pep:*

7: /cgn2 6/ptodata/1/paa/USO3_NEW_COMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-99-602-24

PCT-USO2-21461-3

US-09-809-602-24

US-09-866-99

US-10-274-874-4

US-60-427-552-118

US-60-427-579-118

US-60-427-579-118

US-60-427-579-118

US-60-427-579-118

US-60-427-579-118

US-60-427-579-118

US-09-701-080C-22

US-09-701-080C-22

US-09-701-080C-22

US-09-701-080C-22

US-09-701-080C-22

US-09-701-080C-22

US-09-701-080C-22

US-10-218-286

US-10-218-282-898

US-10-233-553-26

US-10-233-553-28

US-10-233-553-28
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Maximum Match 100%
Listing first 45 summaries
                                                         OM protein - protein search, using sw model
                                                                                                                                                                 105
1 PPLSQETFSDLWKLLPENG 19
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Maximum DB seq length: 200000000
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Match Length DB
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51.5
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3470, Ap
6, Appli
77164, A
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1032, Ap
29, Appl
156, App
2160, Ap
 3137, Ap
4695, Ap
7, Appli
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77164, A
77166, A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 351;
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; Sequence 56463, Application US/09724676A
; Sequence 56463, Application US/09724676A
; GENERAL INFORMATION:
; TITLE OF INVENTION: Variants of alternative splicing
; TITLE OF INVENTION: Variants of alternative splicing
; FILE REPERENCE: 129181.4 Compugen
; CURRENT FILING DATE: 2000-11-28
; SOFTWARE: PATENTING DATE: 2000-11-28
; SOFTWARE: PATENTIN version 3.2
; SOFTWARE: PATENTIN version 3.2
; SEQ IN NO 5463
; TYPE: PAT
; ORCANISM: Homo sapiens
US-09-724-676A-56463
                                                                                                                                                                                                                                                                                                                         APPLICANT: Compugen LTD
TITLE OF INVENTION: Variants of alternative splicing
FILE REFERENCE: 129181.4 Compugen
CURRENT APPLICATION NUMBER: US/09/724,676
CURRENT FILING DATE: 2000-11-28
NUMBER OF SEQ ID NOS: 97222
SEQ TRARE: PatentIn version 3.2
SEQ ID NO 55463
LENGTH: 351
                    PCT-USO2-32851-7
US-10-072-012-607
US-10-072-012-608
US-10-072-012-608
US-10-031-822-49
PCT-USO2-3277-3470
US-10-057-498-3470
US-10-057-498-3470
US-09-724-676-77164
US-09-724-676-77164
US-09-724-676-77166
US-09-724-676-77166
US-10-163-587A-17
US-10-276-781-1032
PCT-USO2-32851-29
PCT-USO2-32851-29
PCT-USO2-32851-29
PCT-USO2-32851-29
US-10-264-237-2160
US-10-264-237-2160
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100.0%; Pred. No. v...
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; Sequence 56463, Application US/09724676
; GENERAL INFORMATION:
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Best Local Similarity 100.
Matches 18, Conservative
 397
434
837
2828
2828
2828
70
70
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1481
1481
1481
1481
1724
1763
612
617
208
                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                 US-09-724-676-56463
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Methods of Use
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PRIOR APPLICATION NUMBER: US/09/277,196
PRIOR FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: 60/079736
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Smith, Timothy S. REGISTRATION NUMBER: 35,367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (415) 576-0300 INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 393 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: Protein
Wills, Ken N.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                       STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
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US-10-274-874-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94.3%; Score 99; DB 5; Length 353; 100.0%; Pred. No. 6.3e-08; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: BIXIL PHARMACCULICALS, INC. et al.
TITLE OF INVENTION: SIR2 ACTIVITY
FILE REPERBENCE: 13407-015W01
CURRENT APPLICATION NUMBER: PCT/US02/21461
CURRENT PILING DATE: 2002-07-08
FRICR APPLICATION NUMBER: US 10/190,159
FRICR APPLICATION NUMBER: US 60/303,370
FRICR APPLICATION NUMBER: US 60/303,456
FRICR FILING DATE: 2001-07-06
F
                                                                                                                                                                                                                                                                                                       APPLICANT: Old, Lloyd J.
APPLICANT: Stockert, Elisabeth
APPLICANT: Stockert, Elisabeth
APPLICANT: Chen, Yao-Tseng
FILE REFERENCE: L0461/7105 (JRV)
CURRENT APPLICATION NUMBER: US/09/849,602
CURRENT PILLING DATE: 2001-05-04
NUMBER OF SEQ ID NOS: 30
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME / KEY: UNSURE
LOCATION: (76)..(76)
OTHER INFORMATION: X = any amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 9, Application US/09860286
GENERAL INFORMATION:
APPLIÉANT: Gregory, Richard J.
                                                                                                                                                                                          Sequence 24, Application US/09849602
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                           APPLICANT: Scanlan, Matthew J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PPLSQETFSDLWKLLPEN 18
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        12 PPLSQETFSDLWKLLPEN 29
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Best Local Similarity 100.0
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
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Best Local Similarity
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LENGTH: 393
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US-09-860-286-9
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ZUP: 94111-3834

COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC COMPAtible
COMPUTER: BAFTEN: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/860,286
FILING DATE: 18 -May-2001
CLASSIFICATION SURKNOWN>
PRIOR APPLICATION DATA:

RELEASTOR TO BATA: COMPANIENT OF THE CONTROL OF THE CONTR
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Maneval, Daniel C. TILLE OF INVENTION: Recombinant Adenoviral Vector and
                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                  STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Trink, Barry
APPLICANT: Ratovitski, Edward
APPLICANT: Sidransky, David
TITLE OF INVENTION: P40 Protein Acts as an Oncogene
FILE REFERENCE: 010.779765
CURRENT APPLICATION NUMBER: US/10/274,874
CURRENT FILING DATE: 2002-10-22
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OTHER INFORMATION: /note= "human p53"
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A APPLICATION NUMBER: US/08/328,673
FILING DATE: 25-OCt-1994
APPLICATION NUMBER: US 08/142,669
FILING DATE: 25-OCT-1993
APPLICATION NUMBER: US 08/233,669
FILING DATE: 26-APR-1994
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APPLICANT: Bright, Gary R.
APPLICANT: Bright, Gary R.
APPLICANT: Premkumar, D. David
APPLICANT: Chen, Yih-Tai
TITLE OF INVENTION: Yih-Tai
FILE REPERENCE: 01-1022-05
CURRENT APPLICATION NUMBER: US/10/211,088
CURRENT PRING DATE: 2002-10-15
PRIOR PEPLICATION NUMBER: 60/309,395
PRIOR FILING DATE: 2001-08-01
PRIOR APPLICATION NUMBER: 60/341,589
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US-60-438-805-9
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; GENERAL INFORMATION:
; APPLICANT: Hyseq, Inc.;
; APPLICANT: Tang, Y, Tom et al
; TITLE OF INVENTION: Novel Nucleic Acids and Polypeptides
; FILE REFERENCE: 21272-030
; CURRENT PELLORION NUMBER: US/10/276,774
; CURRENT FILING DATE: 2002-11-18
; PRIOR FILING DATE: 2000-04-27
; PRIOR FILING DATE: 2000-04-27
; PRIOR FILING DATE: 2000-04-27
; NUMBER OF SEQ ID NOS: 2700
; SOFTWARE: CUSTOM
                                                                                                                                                                                                                                                                                                                                                                                                                                              87.6%; Score 92; DB 7; Length 19;
100.0%; Pred. No. 4.5e-08;
tive 0; Mismatches 0; Indels
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                                         Sequence 9, Application US/60438805
GENERAL INFORMATION:
APPLICANT: KODADEK, THOMAS
TITLE OP INVENTION: MIXED ELEMENT CAPTURE AGENTS
FILE REFERENCE: UTSD: 935USP1
CURRENT APPLICATION NUMBER: US/60/438,805
CURRENT APPLICATION NUMBER: 2003-01-09
NUMBER OF SEQ ID NOS: 12
SOFTWARE: Patentin Ver. 2.1
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                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Artificial Sequence
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Best Local Similarity 100."
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Matches 17; Conservative
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CORGANISM: Homo sapiens
US-10-276-774-2505
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US-10-276-774-2505
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                           US-60-438-805-9
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LENGTH: 19
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING MAJOR DEPRES
TITLE OF INVENTION: DISORDER
TITLE OF INVENTION: DISORDER
TITLE OF DISORDER: US/60/423,552
CURRENT APPLICATION NUMBER: US/60/423,552
CURRENT FILING DATE: 2002-11-05
NUMBER OF SEQ ID NOS: 256
SEQ ID NO 118
LENGTHRE: Patentin version 3.1
SEQ ID NO 118
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TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR DIAGNOSING AND TREATING MAJOR DEPRES
TITLE OF INVENTION: DISORDER
TITLE REPERENCE: AM101250L
CURRENT APPLICATION NUMBER: US/60/427,579
CURRENT APPLICATION NUMBER: US/60/427,579
NUMBER OF SEQ ID NOS: 256
SOFTWARE: Patentin version 3.1
SEQ ID NO 118
LENGTH: 393
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                                         NUMBER OF SEQ ID NOS: 20 SOFTWARE: PastSEQ for Windows Version 3.0
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WS-60-423-552-118
; Sequence 118, Application US/60423552
; GENERAL INFORMATION:
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PRIOR FILING DATE: 1998-03-27
                                                                                                                                                                                                                                                                                                             1 PPLSQETFSDLWKLLPEN 18
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Les 18; Conservative
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                                                                                  ; SEQ ID NO 4
; LENGTH: 393
; TYPE: PRT
; CRGANISM: Homo sapiens
US-10-274-874-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
CORGANISM: Homo sapiens
US-60-423-552-118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
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Matches
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; OTHER INFORMATION: Description of Artificial Sequence:derived from p53
US-09-701-080C-27
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APPLICANT: Shimkets, Richard A.
TITLE OF INVENTION: NOVEL POLYNUCLECTIDES AND POLYPEPTIDES
TITLE OF INVENTION: BNCODED THEREBY
FILE REFERENCE: 15966-543 CON
CURRENT APPLICATION NUMBER: 02/10/218,140
CURRENT PILING DATE: 2002-08-12
PRIOR PELLING DATE: 2000-03-30
PRIOR PLING DATE: 1999-04-05
PRIOR PLING DATE: 1999-04-05
PRIOR PLING DATE: 1999-04-02
PRIOR APPLICATION NUMBER: 60/127,636
PRIOR PLING DATE: 1999-04-02
PRIOR PLING DATE: 1999-04-02
PRIOR PLING DATE: 1999-04-02
PRIOR PLING DATE: 1999-03-11
NUMBER OF SEQ ID NOS: 6322
                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sidney, John
APPLICANT: Sidney, John
APPLICANT: Southwood, Scott
APPLICANT: Belmwund, Inc.
ITILE OF INVENTION: HLA Binding Peptides and Their Uses
FILE REFERENCE: 39963-20019_20
CURRENT FILING DATE: 1998-111-10
PRIOR APPLICATION NUMBER: US/09/189,702A
CURRENT FILING DATE: 1998-111-10
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1998-03-04
NUMBER OF SEQ ID NOS: 380
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 286
LENGTH: 11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 56.2%; Score 59; DB 5; Length 11; Best Local Similarity 100.0%; Pred. No. 0.0046; Matches 11; Conservative 0; Mismatches 0; Indels
                                                                                                    Length 13;
                                                                                                 Query Match 66.7%; Score 70; DB 5; Length 13; Best Local Similarity 100.0%; Pred. No. 9.6e-05; Matches 13; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ), OTHER INFORMATION: p53.14 peptide 34.0324
US-09-189-702A-286
                                                                                                                                                                                                                                                                                                                                                                                    Sequence 286, Application US/09189702A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 250, Application US/10218140; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Sette, Alessandro
APPLICANT: Sidney, John
                                                                                                                                                                                                                                      1 SQETFSDLWKLLP 13
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US-09-189-702A-286
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US-10-218-140-250
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LENGTH: 927
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US-09-701-080C-27
Sequence 27 Application US/09701080C
Sequence 27 Application US/09701080C
GENERAL INFORMATION
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILE REPERRINCE: N73477C GCW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT APPLICATION NUMBER: GB 9811303.8
PRIOR PILING DATE: 1998-05-26
PRIOR FILING DATE: 1998-01-05
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PATCHIN VET: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 22, Application US/09701080C
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: POLYPEPTIDES PROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 F
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILE REPERENCE: N/34/7/C GW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT APPLICATION NUMBER: US/09/701,080C
PRIOR PRILING DATE: 1998-05-26
PRIOR PILING DATE: 1998-05-26
PRIOR APPLICATION NUMBER: GB 9900157.0
PRIOR PILING DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
SOFTWARE: PATENTIN Ver: 2.1
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                                                                                                                                                                                                                                                                                                                                        0; Gaps
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                                                                                                                                                                                                                                                                               Query Match 77.1%; Score 81; DB 6; Length 15; Best Local Similarity 100.0%; Pred. No. 2e-06; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 66.7%; Score 70; DB 5; Length 13; 100.0%; Pred. No. 9.6e-05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                ; OTHER INFORMATION: Binding domain US-10-211-088-141
PRIOR FILING DATE: 2001-12-13
NUMBER OF SEQ ID NOS: 366
SOFTWARE: PatentIn version 3.1
SEQ ID NO 141
                                                                                                                                                        ORGANISM: Artificial sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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Best Local Similarity 100.0
Matches 13; Conservative
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Query Match 49.0%; Score 51.5; DB 6; Length 927; Best Local Similarity 47.4%; Pred. No. 6; Mismatches 3; Indels 1; Gaps Matches 9; Conservative 6; Mismatches 3; Indels 1; Gaps
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1;

Search completed: February 12, 2003, 10:42:49 Job time : 20 secs

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February 12, 2003, 10:35:42; Search time 11 Seconds (without alignments)
44.130 Million cell updates/sec
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                                                              140259 seqs, 25548876 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Maximum DB seq length: 200000000
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                                                                                                                                                                                                                                                       Sequence:
                                                                                                                                                                                                                                                                                                                                                              Searched:
                                                                                                                          Run on:
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Published Applications AA:*

1: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

6: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*

7: /cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*

8: /cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*

9: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*

10: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*

11: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*

11: /cgn2_6/ptodata/1/pubpaa/USO0_PUBCOMB.pep:*

12: /cgn2_6/ptodata/1/pubpaa/USO0_PUBCOMB.pep:*

13: /cgn2_6/ptodata/1/pubpaa/USO0_PUBCOMB.pep:*

14: /cgn2_6/ptodata/1/pubpaa/USO0_PUBCOMB.pep:*

Appl Appli Sequence 74, App. App] Sequence 1 Sequence 1 Sequence 1 Sequence 1 Sequence 1 Sequence 1 Sequence 3 Sequence 3 Sequence 1 Sequence 1 Description Sequence Sequence Sequence Sequence Sequence Sequence Sequence US-10-155-059-8 US-10-155-059-10 US-10-155-059-11 US-10-155-059-16 US-09-776-695-32 US-09-776-695-32 US-09-80-211-9 US-10-155-059-14 US-10-155-059-14 US-10-155-059-13 US-09-214-371-73 US-09-214-371-73 US-09-214-371-73 US-09-214-371-73 US-09-214-371-73 US-09-214-371-73 US-09-214-371-73 US-10-155-059-13 US-10-155-059-13 US-10-155-059-13 US-10-155-059-13 US-09-214-371-74 SUMMARIES Query Match Length DB Score Result

. 0

> ; Sequence 8, Application US/10155059 ; Patent No. US20020147173A1

US-10-155-059-8

RESULT 2

Sequence 18, Appl Sequence 17, Appl Sequence 22, Appl Sequence 39, Appl Sequence 7, Appli Sequence 7, Appli Sequence 24, Appl Sequence 24, Appl Sequence 26, Appl Sequence 25, Appl Sequence 25, Appl Sequence 26, Appl Sequence 26, Appl Sequence 26, Appl	Sequence 34, Appl Sequence 17, Appl Sequence 130, Appl Sequence 112, Appl Sequence 12, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 21, Appl Sequence 2384, Appl Sequence 24, Appl Sequence 24, Appl
12 US-10-155-059-18 10 US-09-214-371-17 10 US-09-950-692-6 10 US-09-155-059-22 10 US-09-214-371-39 9 US-10-024-123-7 10 US-09-214-371-24 10 US-09-214-371-41 10 US-09-214-371-25 10 US-09-214-371-25 10 US-09-214-371-25 10 US-09-214-371-25 10 US-09-214-371-26	10 US-09-214-371-34 10 US-10-155-059-17 10 US-09-0119-679-3 10 US-10-078-770-172 9 US-10-045-792-12 12 US-10-155-059-21 10 US-09-214-371-20 10 US-09-99-630-21 10 US-09-99-630-21 10 US-09-796-692-234 12 US-10-155-059-24 13 US-09-54-433-8
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22 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2	######################################

ALIGNMENTS

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APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Picksley, Stephen
APPLICANT: Picksley, Stephen
APPLICANT: Gracia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
CURRENT FILING DATE: 1999-00-26
PRIOR PELLING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
ASSETWARE: ParentIn Ver. 2.0
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94.3%; Score 99; DB 10; Length 18;
Best Local Similarity 100.0%; Pred. No. 5.5e-09;
Matches 18; Conservative 0; Mismatches 0; Indels
                           ; Sequence 74, Application US/09214371B
; Patent No. US20010018511A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                               Lane, David
                                                                           GENERAL INFORMATION:
APPLICANT: Lane, D
US-09-214-371-74
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LENGTH: 18
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MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-10-155-059-10
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APPLICANT: William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                         TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FESSESO for Windows Version 2.0b
SOFTWARE: PESSESO for Windows Version 2.0b
CURRING APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-AMY-2002
CLASSIFICATION: CUNKNOWN>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Eisenstein, Ronald I
REGISTRATION UNDBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQUENCE DESCRIPTION: SEQUENCE DESCRIPTION: SEQ ID NO: 8:
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                           ADDRESSEE: Nixon Peabody LLP
                                                                                                                                                                                                                   STREET: 101 Federal Street
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Patent No. US20020147173A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                         ZIP: 02110
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617-345-1300
    GENERAL INFORMATION:
APPLICANT: Kaelin, William
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                                                                                                                                                                       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
                                                                                                                                            NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: MA
COUNTRY: USA
ZIP: 02110
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TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEG for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: CUMRNOWN>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Bisenstein, Ronald I
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 11, Application US/10155059; Patent No. US20020147173A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kaelin, William
                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 617-345-1300
TELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
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MUTATIONS WHICH SUPPRESS P53 CANCER MUTA- TIONS
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             94.3%; Score 99; DB 12; Length 20; 100.0%; Pred. No. 6.2e-09; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/776,695
FILING DATE: 26-Feb-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                           Sequence 32, Application US/09776695; Patent No. US20020068283A1; CENERAL INFORMATION:
APPLICANT: Bocke, Jef
Brachmann, Rainer
TITLE OF INVENTION: NUCLEIC ACIDS ENCODING P53
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APPLICANT: Gu, JiJie
TITLE OF INVENTION: Inhibition of p53 Degradation
FILE REPERENCE: 21508-044
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 01107.03170 TELECOMMUNICATION INFORMATION:
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US-09-776-695-32
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FILING DATE: 2001-03-28
ATTORNEY/AGENT INFORMATION:
NAME: Kagan, Sarah A
REGISTRATION NUMBER: 32141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Banner & Witcoff STREET: 1001 G Street, NW CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 393 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: 202-508-9100
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INFORMATION FOR SEQ ID NO: 32:
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                                                Best Local Similarity 100.
Matches 18; Conservative
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Best Local Similarity 100.
Matches 18; Conservative
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US-09-776-695-32
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                    Query Match
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Jost, Christine
Jost, METHONS OF TREATMENT USING
TITLE OF INVENTION: MBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <a href="https://documerrors.org/">CLASSIFICATION</a>: <a href="https://documerrors.org/">ADMINISTRATION</a>: <a hre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTONNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 30,628
REPERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
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SEQUENCE DESCRIPTION: SEQ ID NO: 11:
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REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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Patent No. US20020147173A1
GENERAL INFORMATION:
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                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS: LENGTH: 20 amino acids
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MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: single
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INFORMATION FOR SEQ ID NO: 16:
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STRANDEDNESS: single
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                                                                                                                                                                      'ELEX: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100.
Matches 18; Conservative
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ZIP: 02110
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US-10-155-059-16
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APPLICANT: Gregory, Ren N.
Walls, Ken N.
Maneval, Daniel C.
TITLE OF INVENTION: Recombinant Adenoviral Vector and
Methods of Use
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COMPUTER READABLE FOORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/09/860,211
FILING DATE: 18-MAY-2001
CLASSIFICATION: <a href="https://documents.com/currents-new/">CURRENT APPLICATION</a> (ADMONDA)
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ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, Bighth Floor
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APPLICATION NUMBER: US 08/328,673
FILING DATE: 15-FEB-2000
APPLICATION NUMBER: US 08/142,669
FILING DATE: 25-CT-1993
APPLICATION NUMBER: US 08/233,669
FILING DATE: 26-APR-1994
ATTORNEY/AGENT INFORMATION:
NAME: SMICH, TIMCthy S.
REGISTRATION NUMBER: 35,367
REFERENCE/DOCKET NUMBER: 016930-000920US
TELEPHONE: (415) 576-0300
TELEFAX: (415) 576-0300
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LOCATION: 1..393
ØTHER INFORMATION: /note= "human p53"
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 3
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STRANDEDNESS: <Unknown>
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Patent No. US20020137212A1
GENERAL INFORMATION:
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STATE: California
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                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Homo sapiens
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US-09-860-211-9
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APPLICANT: Kaelin, William
APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
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                                                          94.3%; Score 99; DB 10; Length 393; 100.0%; Pred. No. 1.5e-07; tive 0; Mismatches 0; Indels
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COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATE:
APPLICATION NUMBER: US110/155,059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.3%; Score 99; DB 12; L. 100.0%; Pred. No. 1.5e-07; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: 24 MAY-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
FILING DATE: 12-MAY-1998
FILING DATE: 12-MAY-1997
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
SEQUENCE DESCRIPTION: SEQ ID NO: 9:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-10-155-059-14 ; Sequence 14, Application US/10155059
                                                                                                                                                                                                                                                                                           Sequence 4, Application US/10155059; Patent No. US20020147173A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 394 amino acids
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                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                      12 PPLSQETFSDLWKLLPEN 29
                                                               Query Match
Best Local Similarity 100.0
Matches 18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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Matches 18; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-10-155-059-4
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US-09-860-211-9
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APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Stephen
APPLICANT: Pickeley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Grokeppel, Heinz-Kurt
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INNENTION: Inhibitors of the Interaction of P53 and MDM2
TITLE OF INNENTION: Inhibitors of the Interaction of P53 and MDM2
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT APPLICATION NUMBER: US/09/214,371B
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR APPLICATION NUMBER: PCT/EP97/03549
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 73
LENGTH: 22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 94; DB 12; Length 20; Pred. No. 3.5e-08; 0; Mismatches 1; Indels
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                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
                                                                                                                                                                                                                   PRIOR APPLICATION DATA:

APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REPERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 15:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 73, Application US/09214371B; Patent No. US20010018511A1; GENERAL INFORMATION:
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; OTHER INFORMATION: X = Biotin-Ser
US-09-214-371-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 15:
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ORGANISM: Artificial Sequence
                   COMPUTER READABLE FORM
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Best Local Similarity 94.47
Matches 17; Conservative
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Atent No. Uszuwerton:
GENERAL INFORMATION:
APPLICANT: Kaalin, William
JOSE, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 15, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
APPLICANT: Kaelin, William
Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
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                                                                                                                                                                                                                                                                                                                                                                                          CURRENT IFFE: DISKRETE
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-May-1996
FILING DATE: 12-May-1997
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-May-1997
APPLICATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELEFRANION INFORMATION:
TELEFRAN: 617-345-6054
TELEFRAN: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 14:
US-10-155-059-14
                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 20 amino acids
                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
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ZIP: 02110
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COUNTRY: USA
ZIP: 02110
                                                                                                                                                                                                                                                                Boston
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Matches 17; Conserva
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ATTORNEY/AGENT INFORMATION:
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SOFTWARE: Pat
SEQ ID NO 1
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APPLICANT:
APPLICANT:
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TITLE OF INVENTION: MES-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 19;
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                        0; Indels
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SOFTWARE: FASTEM: Windows
SOFTWARE: FASTEM: Windows
SOFTWARE: FASTEM: Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24-May-2002
CLASSIFICATION - CURROWN>
PRIOR CLASSIFICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
*FILING DATE: 12-MAY-1997
                                                                                                                                                                                                                                                                    APPLICANT: Yean, Zhi-Min
APPLICANT: Yean, Zhi-Min
APPLICANT: Gu, Jijle
TITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION WUMBER: 18/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR FILING DATE: 1999-12-08
PRIOR FILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: PatentIn Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 87.6%; Score 92; DB 10; L Best Local Similarity 100.0%; Pred. No. 6.6e-08; Matches 17; Conservative 0; Mismatches 0;
      Pred. No. 5.5e-08;
                          0; Mismatches
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ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
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Patent No. US20020147173A1
GENERAL INFORMATION:
                                                                                                                                                                                                          ; Sequence 7, Application US/09732384; Patent No. US20020132977A1; GENERAL INFORMATION:
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      Best Local Similarity 100.0%;
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                                                                                             6 PPLSQETFSDLWKLLPE 22
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                          17; Conservative
                                                                  1 PPLSQETFSDLWKLLPE 17
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US-09-732-384-7
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                            Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 84.8%; Score 89; DB 10; Length 19; 94.1%; Pred. No. 1.9e-07; trive 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                              85.7%; Score 90; DB 12; Length 19
88.9%; Pred. No. 1.3e-07;
tive 1; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Dickeley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interac
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT PILING DATE: 1999-03-26
PRIOR PILING DATE: 1997-07-04
NAME: Elsentein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
TELEFATA: «Unknown»
INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Search completed: February 12, 2003, 10:43:07
                                                                                                                                                                                                                                                               TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Lane, David
Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
Hochkeppel, Heinz-Kurt
Garcia-Echeverria, Carlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 1, Application US/09214371B; Patent No. US20010018511A1; GENERAL INFORMATION:
                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
LENGTH: 19 amino acids
                                                                                                                                                                                                                                        STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT ORGANISM: Artificial Sequence
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SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                     1 PPLSQETFSDLWKLLPEN 18
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Best Local Similarity 94.19
Matches 16; Conservative
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                                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 88.99
Matches 16; Conservative
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0

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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 10:32:16; Search time 36 Seconds (without alignments) 70.327 Million cell updates/sec Run on:

Title: US-09-403-440A-2
Perfect score: 105
Sequence: 1 PPLSQETFSDLWKLLPENG 19

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

908470 segs, 133250620 residues Searched:

908470 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	p53 homologue TIP p53 homologue TIP p53 N-terminal pep Immunodominant epi Human p53 amino ac p53 peptide used t Peptide fragment o p53 protein amino p53 protein amino p53 protein amino
SUMMARIES	AAW82321 AAW82319 AAW37228 AAR54907 AAR51879 AAW60202 AAW30523 AAW13603 AAW13603
08	19 19 11 11 11 11 11 11 11 11 11 11 11 1
* Query Match Length DB	119 119 125 126 137 136 136 137 137 137 137 137 137 137 137 137 137
* Query Match	100.0 100.0 94.3 94.3 94.3 94.3 94.3
Score	100 100 000 000 000 000 000 000
Result No.	1008432

Disclosure, Fig 1, 52pp; English.

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14 AAR42174	7 AAW0788	9 AAW5724	19 AAW48243	9 AAW42	9 AAW4297	0 AAW9430	8 AAW4707	2 AAB3668	a	22 AAB36685	a	ď	N	1 AAY7071	2 ABG0151	80	8	8 AAW13960		8 AAW13	8 AAW13	8 AAW139	B AAW1397	8 AAW1397	8 AAW1397	8 AAW1397	80	80	8 AAW1395	8 AAW1395	8 AAW1395	AAR222	L L	AAR9462
64	64		4	4	4	64	_	_	₹#	_	10	_	_	_	~	_		359				363 1	_		363	_				368 1		393 1	393 1	393 1
	σ'n	94.		6		0	94	94.3	94		94.	94.	94.	94.	94.	94.	94.	94.	94.	94.	94.	94.	94.	94	94.	94.	94.	94.	94.3	94	94.	94.		94.
66	66	66	66	66	66	66	66	99	66	66	99	66	66	66	66	99	99	99	66	66	66	66	66	66	66	66	66	66	66	66	66	66	66	66
11	12	13	14	15	16	17	18	19	20	21	22	23	24	25	56	27	28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

AAW82321 standard; Peptide; 19 AA. XX AAW82321; XX AAW82321; XX AAW82321; XX XX XX XX XX YOU DF 22-FEB-1999 (first entry) XX

medicament.

X8888888888888X8

This sequence is a peptide homologue of a region of p53 which binds to mdm2. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in a population of cells. This agent is also used in production of a therapeutic for activating p53, where the population of cells do not overexpress mdm2. Inhibiting mdm2 production and/or inhibiting the clearance of p53 by income of p53 to increase by function. The agents for use in therapeutics for activate p53 function. The agents for use in therapeutics for activate p53 used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.

19 AA; Sequence

0; Gaps 100.0%; Score 105; DB 19; Length 19; 100.0%; Pred. No. 5.7e-10; ive 0; Mismatches 0; Indels C 1 PPLSQETFSDLWKLLPENG 19 1 PPLSOETFSDLWKLLPENG 19 Best Local Similarity 100.0 Matches 19; Conservative Query Match ò a

. 0

RESULT 2

AAW82319 standard; Peptide; 19 AA. AAW82319

AAW82319;

22-FEB-1999 (first entry)

p53 homologue TIP peptide.

p53; mdm2; inhibitor; therapy; activator; treatment; cancer; medicament.

Synthetic.

W09847525-A1.

29-OCT-1998.

Compounds binding to MDM2 protein and inhibit its interaction with p53 - useful in, e.g. diagnosis and treatment of cancer and viral infections and identifying binding agents

Disclosure; Page 34; 45pp; English.

98WO-GB01144. 20-APR-1998;

97GB-0008092. 22-APR-1997;

(UYDU-) UNIV DUNDEE.

Lane DP;

WPI; 1998-609932/51.

New agents which inhibit interaction of p53 and mdm2 - useful for activating p53, e.g. for treating cancers, viral conditions or other conditions associated with non functional p53 or mdm2

Disclosure, Fig 1; 52pp; English.

This sequence is a peptide homologue of a region of p53 which binds to made. This peptide is used in the construction of a novel agent capable of disrupting the binding of p53 and mdm2 or inhibiting the production of mdm2 in apoulation of cells. This agent is also used in the preparation of a therapeutic for activating p53, where the population of cells do not overexpess mdm2. Inhibiting mdm2 production and/or inhibiting the binding of mdm2 to p53 allows levels of p53 to increase by inhibiting the clearance of p53 by mdm2, and can be used to activate p53 function. The agents for use in therapeutics for activating p53 can be used for the treatment of cancer, viral conditions or other conditions associated with non-functional p53.

19 AA; Sequence

Query Match

100.0%; Score 105; DB 19; Length 19;

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0
                                                                                                                                MDM2; oncogenic protein; p53; human; inhibition; interaction; cancer; tumour; diagnosis; binding; viral infection; Elisa TIP assay.
          Gaps
                                                                                                                                                                                                                                                      Boettger V, Chene P, Furet P, Garcia-echeverria C;
Lane D, Picksley S;
          ..
           Indels
                                                                                                                   p53 N-terminal peptide fragment for Elisa TIP assay.
           ..
    5.7e-10;
    Pred. No. 5.7
, Mismatches
                                                                                                                                                                                                                                    (CANC-) CANCER RES CAMPAIGN TECHNOLOGY.
                                                                           AAW37228 standard; peptide; 18 AA.
97WO-EP03549.
                                                                                                                                                                                                                97GB-0007041.
                                                                                                                                                                                                                      96GB-0014197.
                          1 PPLSQETFSDLWKLLPENG 19
                                   1 PPLSQETFSDLWKLLPENG 19
                                                                                                        (first entry)
            Conservative
                                                                                                                                                                                                                                                                 Hochkeppel H, Lane D,
                                                                                                                                                                                                                                            (NOVS ) NOVARTIS AG.
                                                                                                                                                                                                                                                                               WPI; 1998-100996/09.
      Best Local Similarity
                                                                                                                                                          Homo sapiens.
                                                                                                                                                                       WO9801467-A2.
                                                                                                                                                                                                                  07-APR-1997;
                                                                                                                                                                                                                        05-JUL-1996;
                                                                                                        20-JUL-1998
                                                                                                                                                                                                                                                           Boettger A,
                                                                                                                                                                                     15-JAN-1998.
               19;
                                                                                          AAW37228;
              Matches
                                                                                          δ
                                         Ωp
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This represents a p53 N-terminal peptide fragment used in an Elisa TIP assay for analysing the interaction between human encogenic protein MDM2 assay for analysing the interaction between human encogenic protein MDM2 and p53. The invention provides peptide derivatives capable of binding to the human MDM2 These peptides can specifically inhibit or block the binding of MDM2 to the human p53 protein, in vitro or in vivo. Inhibiting the interaction between the p53 and MDM2 can induce growth arrest or apoptosis in tumour cells comprising a wild-type p53 and non-elevated composits in tumour cells our provision and induce that bind levels of MDM2. The peptides may be used to identify and design inhibitors of MDM2, p53 binding. They to MDM2 and to identify and design inhibitors of MDM2, p53 binding. They compared the poptides of MDM2, and leukaemia classase how measuring levels of MDM2 in blood of cancer and leukaemia chiesease involving p53/MDM2 contents and for treatment or prevention of disease involving p53/MDM2 contents and for treatment or prevention of disease involving p53/MDM2 contents and interactions. Phe peptides can administered nasally, rectally, orally or by injection. By interfering with MDM2/p53 interaction, the peptides can accivate p53 function and accumulation in normal cells. The peptides which mind; the MDM2 banding content with wild-type p53. <u>`</u> Gaps ; Length 18; 0; Indels 94.3%; Score 99; DB 19; L 100.0%; Pred. No. 4.9e-09; iive 0; Mismatches 0; Query Match Best Local Similarity 100.0

1 PPLSQETFSDLWKLLPEN 18

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Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptides derived from the N-terminal (amino acids 1-112) or the C-terminal (amino acids 350-393) of protein p53 which specifically react with anti-p53 antibodies in patients with cancer or precancerous conditions are claimed. The peptides (AAR54907-R54921) are useful for detecting and monitoring cancerous and precancerous
                                                                                                                                                                 cancer; pre-cancerous state; detection; diagnosis; human p53 gene;
immunodominant epitope; human cellular tumour antigen;
transformation-associated protein.
                                                                                                                                                                                                                                                                                                                                                                                                                         New immuno:dominant epitope(s) of protein p53 - for detecting and monitoring antibodies indicative of cancer and precancerous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human nuclear phosphoprotein p53; tumour suppressor gene product; anti-oncogene; cancer; tumour; antibody binding region; epitope.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94.3%; Score 99; DB 15; Length 25; 100.0%; Pred. No. 7.1e-09; ive 0; Mismatches 0; Indels
                                                                                                                                      Immunodominant epitope from p53 N-terminal.
                                                              AAR54907 standard; peptide; 25 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AARS1879 standard; Protein; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 4; Page 42; 62pp; French.
                                                                                                                                                                                                                                                                                                                                                                            Soussi T;
                                                                                                                                                                                                                                                                                                                         92FR-0013110.
                                                                                                                                                                                                                                                                                                93WO-FR01082.
1 PPLSQETFSDLWKLLPEN 18
                                                                                                                (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 PPLSQETFSDLWKLLPEN 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human p53 amino acids 9-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-NOV-1994 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                (EURO-) LAB EUROBIO SA
                                                                                                                                                                                                                                                                                                                                                                        Legros Y, Lubin R,
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1994-167463/20.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            25 AA;
                                                                                                                                                                                                                                                                                             02-NOV-1993;
                                                                                                                                                                                                                                                                                                                       02-NOV-1992;
                                                                                                              29-NOV-1994
                                                                                                                                                                                                                                            WO9410306-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                     11-MAY-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   conditions.
                                                                                      AAR54907;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR51879;
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                                     RESULT 4
                                                  AARS4907
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 5
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Gaps
                                                                                                                                                                                                                                                                Antibodies specific for p53 are detected by binding to immobilised
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                                                                                                                                                                                                                                                                             fragments of the p53 gene product containing the antibody-binding region. Preferred fragments contain amino acids 1-241, 40-349, 40-383, 66-241, 66-393, 237-349, 237-393 and esp. 9-33, 37-52 or 368-386. See AAR51872-R51881 for sequences of these fragments.
                                                                                                                                                                          Non-radioactive detection of p53 specific antibodies - by capture on immobilised p53 or its fragments, then reaction with labelled second antibody, for diagnosis of tumours and suitable for
                                                                                                                                                                                                                                                                                                                                                                                                  0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; p53; antibody; detection; biosensor; cancer patient;
p53 gene therapy; immune response; mutant.
                                                                                                                   Zentgraf H;
                                                                                                                                                                                                                                                                                                                                                                     94.3%; Score 99; DB 15; Length 25; 100.0%; Pred. No. 7.1e-09; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       p53 peptide used to detect antibodies against p53
                                                                                                                Volkmann M,
                                                                                    (DEKR-) DEUT KREBSFORSCHUNGSZENTRUM STIFTUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW60202 standard; peptide; 26 AA.
                                                                                                                Tessmer C,
                                                                                                                                                                                                                                       Claim 11; Page 19; 35pp; German.
                                      93WO-EP02666
                                                               92DE-4232823
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97WO-US16132.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    96US-0028533.
                                                                                                                                                                                                                                                                                                                                                                                                                                       4 PPLSQETFSDLWKLLPEN 21
                                                                                                                                                                                                                                                                                                                                                                                                                      1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0
Matches 18, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-AUG-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Swanson SJ;
                                                                                                             Klein R, Schranz P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (SCHE ) SCHERING CORP
                                                                                                                                    WPI; 1994-135732/16.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 1998-240965/21.
                                                                                                                                                  N-PSDB; AAQ62365.
                                                                                                                                                                                                                                                                                                                                              25 AA;
                                     30-SEP-1993;
                                                              30-SEP-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Homo sapiens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-OCT-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   07-OCT-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9815834-A1
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                                                                                                                                                                                                                   screening
                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW60202;
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WO9408241-A.

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Peptides AAW60202-05 are derived from human p53 protein. The present peptide corresponds to residues 11-35. The peptides are used in the method of the invention. Antibodies that bind to p53 protein are detected by immobilising a p53 peptide directly on to a flow cell of patient serum, diluted in buffer, and measuring binding of antibody to the peptide using the biosensor. The method is used to monitor cancer patients undergoing p53 gene therapy (to determine if an immune response has developed), and also to detect antibodies against mutant forms of p53. Claim 4; Page 5; 41pp; English. Sequence 8X55555555555X8

26 AA;

0; Gaps Score 99; DB 19; Length 26; 0; Indels 100.0%; Pred. No. 7.4e-09; cive 0; Mismatches 0; 94.3%; 1 PPLSQETFSDLWKLLPEN 18 3 PPLSQETFSDLWKLLPEN 20 18; Conservative Sest Local Similarity Query Match Matches ઠે 셤

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RESULT 7

AAB30523 standard; peptide; 32 AA.

AAB30523;

06-MAR-2001 (first entry)

Peptide fragment of a tumour suppressor protein p53.

Biologically active compound; cellular metabolism; DNA replication; RNA transcription; RNA translation; RNA elongation; RNA processing; protein synthesis; protein processing; cellular differentiation; cell division; ion channel transmission; cellular protein; toxin; RNA transportation; cellular oxidation; tumour suppressor p53; plasminogen antigen activator.

Synthetic.

WO200061775-A1.

19-OCT-2000.

99WO-IB00616. 08-APR-1999; 99WO-IB00616. 08-APR-1999;

(SERG/) SERGEEV P.

Sergeev P;

WPI; 2001-006911/01.

Novel methods for the synthesis of biologically active compounds from inactive precursors in the cells of living organisms, useful for producing proteins or polynucleotides -

Example 8; Page 29; 65pp; English.

The specification describes a method of synthesis of biologically active substances of determined structure directly in the cells of living organisms containing specific RNA or DNA sequence. The method is based on the hybridisation of two or more oligomers bound with biologically inactive substances to specific RNA or DNA in vivo in the cells of living organisms. After hybridisation of the oligomers, the biologically inactive precursors bound to the oligomers can interact with each other to make the active form of the substances. This changing of properties is due to chemical reactions which bind the biologically inactive precursors through a chemical bond into a biologically active form of

active compounds from inactive precursors. These compounds may be active compounds from inactive precursors. These compounds may be transcription, RNA translation, RNA elongation RNA processing, protein synthesis, protein processing, cellular differentiation, cell division, synthesis, protein processing, cellular protein and RNA transportation, cellular protein and RNA transportation, processes of cellular oxidation, cellular protein and RNA transportation, processes of cellular oxidation, cellular protein and RNA transportation, processes of cellular oxidation, cellular protein and RNA transportation, cellular proteins or RNAs. AAB3023-36 represent peptides which are bound to oligomers AAC62167-80 and AAC62181-94. The peptides are fragments of the tumour suppressor p53, and the oligomers are antiparallel to human plasminogen antipan activator mRNA (AAC62167-80) or human NDHOT (AAC62181-94). The method of the invention is used to produce the tumour suppressor protein p53 from the bound peptides and oligomers. the whole compound. The methods are useful for producing biologically 32 AA; Sequence

Gaps ; 0 Length 32; 0; Indels 94.3%; Score 99; DB 22; L 100.0%; Pred. No. 9.2e-09; iive 0; Mismatches 0; 18; Conservative Local Similarity Query Match Matches

1 PPLSQETFSDLWKLLPEN 18 à 12 PPLSQETFSDLWKLLPEN 29

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RESULT 8

AAW13604 standard; peptide; 36 AA. AAW13604

AAW13604;

16-JAN-1998 (first entry)

p53 protein amino acids 6-41.

Mouse, Mdm2, murine double minute; phosphoprotein; binding; modulation; tumour suppressor; p53; oncogene; cell cycle arrest; p107; antagonist; inhibition; transcription factor; adenocarcinoma; colon; cancer; breast; lung; scomach; myeloid leukaemia; lymphoma; hyperproliferative;

Homo sapiens.

restenosis.

WO9709343-A2

13-MAR-1997.

96WO-FR01340. 02-SEP-1996; 95FR-0010331. 04-SEP-1995;

(INRM) INST NAT SANTE & RECH MEDICALE.

(RHON) RHONE POULENC RORER SA.

WPI; 1997-192837/17.

Dubs-Poterszman M, Tocque B,

Wasylyk B;

Treating cancer with antagonist of oncogenic activity of protein Mdm2 - or nucleic acid encoding an antagonist, also viral vectors contg. this nucleic acid

Claim 4; Page -; 43pp; French.

wild type human p53 protein. This peptide corresponds to amino acids 6-41 of the p53 sequence. The peptides are claimed peptides which are able to bind the N-terminal amino acids (1-134) of the murino double minute-2 (mdm2) protein (AAM13600). Mdm2 protein is a 90 KD phosphoprotein which binds and modulates the activity of the tumour suppressor protein p53. It has now been shown that the mdm2 protein itself has oncogenic properties, especially in a p53-mul background. Mdm2 is observed to unblock cell cycle arrest in G1 caused by The peptides AAW13602-6 represent peptide fragments derived from the

constructed from the wild type human p53 sequence.

41 AA;

Sequence

SXS

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over-expression of the p107 protein. The p53 peptides are examples of antagonists of the invention which are able to inhibit the oncogenic activity of mdm2. The antagonists are used to treat e.g. adenocarcinoma of the colon; cancer of the breast, lung or stomach; myeloid leukaemia; Ecall lymphoma, or other hyperpoliferative conditions such as restenosis. Note: this sequence is not given in the specification but is constructed from the wild type human p53 sequence.
                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                  0
                                                                                                                                                                                                                     Length 36;
                                                                                                                                                                                                               Score 99; DB 18; Length 36; Pred. No. 1.1e-08; 0; Mismatches 0; Indels
                                                                                                                                                                                                                 94.3%; S
100.0%;
                                                                                                                                                                                                                                                         18; Conservative
                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                       36 AA;
                                                                                                                                                                       Sequence
                                                                                                                                                                                                               Query Match
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88888888888
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1 PPLSQETFSDLWKLLPEN 18 7 PPLSQETFSDLWKLLPEN 24 RESULT ð

AAW13603 standard; peptide; 41 AA. p53 protein amino acids 1-41. 16-JAN-1998 (first entry) AAW13603; AAW13603

Mouse; Mdm2; murine double minute; phosphoprotein; binding; modulation; tumour suppressor; p53; oncogene; cell cycle arrest; p107; antagonist; inhibition; transcription factor; adenocarcinoma; colon; cancer; breast; lung; stomach; myeloid leukaemia; lymphoma; hyperproliferative; WO9709343-A2 Homo sapiens 13-MAR-1997 restenosis.

(INRM) INST NAT SANTE & RECH MEDICALE RHON) RHONE POULENC RORER SA Dubs-Poterszman M, Tocque B, 95FR-0010331. 02-SEP-1996; 04-SEP-1995;

Wasylyk B;

WPI; 1997-192837/17.

Claim 4; Page -; 43pp; French.

Claim 4; Page -; 43pp; French.

restenosis. Note: this sequence is not given in the specification but is itself has oncogenic properties, especially in a p53-null background. Mdm2 is observed to unblock cell cycle arrest in G1 caused by over-expression of the p107 protein. The p53 peptides are examples of antagonists of the invention which are able to inhibit the oncogenic activity of mdm2. The antagonists are used to treat e.g. adenocarcinoma of the colon; cancer of the breast, lung or stomach; myeloid leukaemia; cell lymphoma, or other hyperproliferative conditions such as wild type human p53 protein. This peptide corresponds to amino acids 1-41 of the p53 sequence. The peptides are claimed peptides which are able to bind the N-terminal amino acids (1-134) of the murine double minute-2 (mdm2) protein (AMM13600). Mdm2 protein is a 90 kD phosphoprotein which binds and modulates the activity of the tumour suppressor protein p53. It has now been shown that the mdm2 protein peptides AAW13602-6 represent peptide fragments derived from the

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                                                                                                                                                                                                tumour suppressor; p53; oncogene; cell cycle arrest; p107; antagonist; inhibition; transcription factor; adenocarcinoma; colon; cancer; breast; lung; stomach; myeloid leukaemia; lymphoma; hyperproliferative;
                                                                                                                                                                                    Mouse; Mdm2; murine double minute; phosphoprotein; binding; modulation;
                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                  Treating cancer with antagonist of oncogenic activity of protein Mdm2 - or nucleic acid encoding an antagonist, also viral vectors contg. this nucleic acid
                     ;
0
   Length 41;
                    0; Indels
 Score 99; DB 18; I
Pred. No. 1.2e-08;
                    Mismatches
                                                                                                                                                                                                                                                                                                                                                (INRM ) INST NAT SANTE & RECH MEDICALE.
                                                                                                         AAW13602 standard; peptide; 52 AA.
         100.0%; Pr
                                                                                                                                                                                                                                                                                                                                                           (RHON ) RHONE POULENC RORER SA.
                                                                                                                                                                                                                                                                                                                                                                             Dubs-Poterszman M, Tocque B,
                                                                                                                                                                  p53 protein amino acids 1-52.
 94.3%;
                                                                                                                                                                                                                                                                                                           96WO-FR01340.
                                                                                                                                                                                                                                                                                                                               95FR-0010331.
                                     1 PPLSQETFSDLWKLLPEN 18
                                                 12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                 (first entry)
                  18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                 WPI; 1997-192837/17.
         Best Local Similarity
                                                                                                                                                                                                                                                Homo sapiens.
                                                                                                                                                                                                                                                                   WO9709343-A2.
                                                                                                                                                16-JAN-1998
                                                                                                                                                                                                                                                                                                          02-SEP-1996;
                                                                                                                                                                                                                                                                                                                              04-SEP-1995;
                                                                                                                                                                                                                                                                                      13-MAR-1997.
                                                                                                                                                                                                                              restenosis.
                                                                                                                            AAW13602;
Query Match
                  Matches
                                                                                       RESULT 10
                                                                                               AAW13602
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The peptides AAW13602-6 represent peptide fragments derived from the wild type human p53 protein. This peptide corresponds to amino acids 1-52 of the p53 sequence. The peptides are claimed peptides which are able to bind the N-terminal amino acids (1-134) of the murine double computed. (mdm2) protein (AAW1360). Mdm2 protein is a 90 kD phosphoprotein which binds and modulates the activity of the tumour cruppressor protein p53. It has now been shown that the mdm2 protein crublors protein crublock cell cycle arrest in G1 caused by cover-expression of the p107 protein. The p53 peptides are examples of antagonists of the invention which are able to inhibit the oncogenic cutivity of mdm2. The antagonists are used to treat e.g. adenocarcinoma of the north are able to inhibit the oncogenic activity of mdm2. The antagonists are used to treat e.g. adenocarcinoma of the breast, lung or stomach; myeloid leukaemia; B cell lymphoma, or other hyperproliferative conditions such as constructed from the wild type human p53 sequence.

52 AA; Sequence

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94.3%; Score 99; DB 18; Length 52; 100.0%; Pred. No. 1.6e-08; Best Local Similarity Query Match

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28-JAN-1997 (first entry)

AAW07886;

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This sequence represents the product of the p53 gene. p53 is a tumour suppressor gene and this protein appears to be a member of a group of proteins which regulate normal cellular proliferation and suppression of cellular transformation. Inactivation of the p53 gene has been implicated in the formation, or progression of a wide variety of carcinoma. p53 is thought to interact with the protein product of a gene which is amplified in a number of human tumours, designated MDM2. Polypeptides containing at least amino acids 13-41 of p53, or the DNA encoding these, may be used to inhibit the growth of tumour cells containing MDM2 gene amplification.
                                                                                                                                                                                p53 gene; tumour suppressor gene; regulation; cellular proliferation; cellular transformation; carcinoma; human; tumour; MDM2; inhibition;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                Diagnosing neoplasia from amplification of MDM2 gene - or elevated gene expression, also new DNA, MDM2 protein, antibodies and treatment of sarcoma by inhibiting MDM2 expression
  ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94.3%; Score 99; DB 14; Length 64; 100.0%; Pred. No. 2e-08; ive 0; Mismatches 0; Indels
  0; Indels
                                                                                                                                                                                                                                                                     /note= "Site of interaction with MDM-2"
                                                                                                                                                                                                                                                                                                                                                                                                            Kinzler KW, Vogelstein B;
   Mismatches
                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 35; 75pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW07886 standard; protein; 64 AA.
                                                                                                 AAR42174 standard; peptide; 64 AA.
     .;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          94.3%;
                                                                                                                                                                                                                                                                                                                                                          92US-0867840
                                                                                                                                                                                                                                                                                                                                    93WO-US03199
                                                                                                                                                                                                                                                                                                                                                                    92US-0903103
                                                                                                                                                                                                                                                                                                                                                                                       (UYJO ) UNIV JOHNS HOPKINS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              12 PPLSQETFSDLWKLLPEN 29
                       1 PPLSOETFSDLWKLLPEN 18
                                   12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                          05-MAY-1994 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18; Conservative
    Conservative
                                                                                                                                                               p53 N-terminal fragment.
                                                                                                                                                                                                                                                                                                                                                                                                            Burrell M, Hill DE,
                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1993-336944/42.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                         gene amplification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      64 AA;
                                                                                                                                                                                                                                                                                                                                      07-APR-1993;
                                                                                                                                                                                                                                                                                                                                                          07-APR-1992;
                                                                                                                                                                                                                                                             Binding-site
                                                                                                                                                                                                                               Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                     23-JUN-1992;
                                                                                                                                                                                                                                                                                                               14-OCT-1993.
                                                                                                                                                                                                                                                                                           WO9320238-A
     18;
                                                                                                                       AAR42174;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 12
AAW07886
ID AAW078
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
     Matches
                                                                             RESULT 11
                                                                                         AAR42174
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AAW07886 represents the human p53 protein which is involved in the development of many cancers. The protein is used here in a method for identifying compounds that interfere with the binding of p53 and NDM-2. In binding the p53 protein, the MDM-2 protein releases a cell from p53-regulated growth, allowing cancers to develop. Therefore compounds identified as interfering with the binding of MDM-2 to p53 are potentially useful in the treatment of human neoplastic cells. In the method pref. one or both of the proteins is a fusion protein esp. with an antibody or antibody fragment which aids separation and identification.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Identification of cpds. interfering with human MDMZ/p53 binding useful as therapeutic agents to treat human neoplastic cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0;
                                              p53; MDM-2; binding-inhibitor; identification; tumour; cancer; neoplasia; antibody fusion protein; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94.3%; Score 99; DB 17; Length 64; 100.0%; Pred. No. 2e-08; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                              13..57
/note= "MDM-2 binding fragment"
                                                                                                                                                                                                                                                                                   /note= "MDM-2 binding fragment"
                                                                                                                                                                                                                                     /note= "MDM-2 binding fragment"
Human p53, involved in tumour suppression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 16; Column 19-20; 36pp; English.
                                                                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAW57240 standard; protein; 64 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human p53 N-terminal fragment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    92US-0867840.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 92US-0903103.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             92US-0867840.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93US-0044619
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94US-0245500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYJO ) UNIV JOHNS HOPKINS.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           12 PPLSOETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                10-AUG-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Kinzler KW, Vogelstein B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-401591/40.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
nes 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            64 AA;
                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               07-APR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             18-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         07-APR-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                             US5550023-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                              27-AUG-1996.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW57240;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                 Region
                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                      Region
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW57240
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The present sequence represents a human p53 fragment 13-41 which is necessary for the interaction of MDM-2 and p53. The present invention describes a method for diagnosing a neoplastic disease caused by overexpression of MDM2 protein. The method comprises detecting an elevated cellular amount of this protein. The method is useful for the diagnosis of sarcoma, especially liposarcoma, malignant fibrous histiocytoma or osteosarcoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human MDM2 binding polypeptide - comprises fragments of p53, useful in re-establishing p53-regulated growth control in cells
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MDM2; tumour; diagnosis; neoplasia; DNA binding protein; p53 polypeptide; binding; tumour cell; p53-regulated growth; inhibition; anti-cancer agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 99; DB 19; Length 64;
Pred. No. 2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                      Cancer diagnosis - by determination of MDM2 protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vogelstein B;
                                                                                                                                                                                                              Burrell M, Hill DE, Kinzler KW, Vogelstein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             N-terminal region of p53 (amino acids 13-41).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                                                                                                                                                                                                                                                                             Disclosure; Column 19-20; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94.3%; Scur
100.0%; Pre
0; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW42878 standard; Protein; 64 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Kinzler KW,
                                                      95US-0390517
                                                                                          93US-0044619
                                                                                                         92US-0867840.
                                                                                                                                              95US-0390517
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  95US-0390516.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93US-0044619.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    92US-0867840.
                                                                                                                                                                              (UYJO) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (UYJO ) UNIV JOHNS HOPKINS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hill DE,
                                                                                                                                                                                                                                                      WPI; 1998-239206/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 18; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-100408/09,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              64 AA;
                                                      17-FEB-1995;
                                                                                                         07-APR-1992;
23-JUN-1992;
                                                                                        07-APR-1993;
                                                                                                                                              17-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-FEB-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         30-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     07-APR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    07-APR-1992;
23-JUN-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US5708136-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            13-JAN-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Burrell M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW42878;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW42878
셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        The method has been developed for inhibiting the growth of tumour cells containing a human MDM2 gene amplification. The method comprises treating the tumour cells with a DNA molecule that expresses a polypeptide capable of binding to human MDM2 protein. The present comprises a polypeptide capable of binding to human MDM2 protein. The present capable of present invention describes three preferred polypeptides for binding human MDM2: (1) the polypeptide comprises amino acids 1-50 of p53 (see AAW7240); (2) the polypeptide comprises amino acids 13-41 of p53 (see AAW7240) and at least none additional p53 residues on the N- or C-terminal side, provided that the polypeptide comprises amino acids 13-41 of p53 (see AAW57241) and at least nine comprises amino acids 13-41 of p53 (see AAW57241) and at least nine comprises amino acids 13-41 of p53 (see AAW57241) and at least nine polypeptide lacks amino acids 13-93 of p53; Some malignant fibrous histocytomas and liposarcomas have an MDM2 gene amplification, so detection of increased expression of MDM2 gene products indicates
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Human; MDM2; hMDM2; tumour; cancer; diagnosis; neoplastic disease; p53; sarcoma; liposarcoma; malignant fibrous histiocytoma; osteosarcoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                         Inhibiting growth of tumour cells having MDM2 gene amplification with p53 protein fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ·,
       Human; p53; MDM2; tumour; growth inhibition; amplification;
malignant fibrous histiocytoma; liposarcoma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   94.3%; Score 99; DB 19; Length 64; 100.0%; Pred. No. 2e-08; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 1; Column 19-20; 40pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAW48243 standard; peptide; 64 AA.
                                                                                                                                                                                                       93US-0044619.
92US-0867840.
92US-0903103.
95US-0390515.
                                                                                                                                                                       95US-0390515.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94.3%;
                                                                                                                                                                                                                                                                                             SNING OUND ( OCYD)
                                                                                                                                                                                                                                                                                                                                       Vogelstein B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            12 PPLSQETFSDLWKLLPEN 29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human p53 fragment 13-41.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18; Conservative
                                                                                                                                                                                                                                                                                                                                                                       WPI; 1998-321574/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   64 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tumourigenesis.
                                                              Homo sapiens
                                                                                                                                                                       17-FEB-1995;
                                                                                              US5756455-A.
                                                                                                                                                                                                         07-APR-1993;
                                                                                                                                                                                                                                                                                                                                 Kinzler KW,
                                                                                                                                                                                                                            07-APR-1992;
                                                                                                                                                                                                                                                            17-FEB-1995;
                                                                                                                                                                                                                                              23-JUN-1992
                                                                                                                                   26-MAY-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
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Gaps

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us-09-403-440a-2.open.rag

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over-expressing MDM2

Claim 1; Columns 19-20; 41pp; English.

The present sequence represents a N-terminal portion of the p53 protein. These amino acid residues have been found to be necessary for the interaction of MDM2 and p53. A cell containing three recombinant but constructs was produced. These constructs encode an MDM2 protein fused to a sequence-specific DNA binding domain, a p53 polypeptide fused to a transcriptional activation domain, and a reporter gene downstream from a DNA element which is recognised by the sequence-specific DNA-binding domain. The cell is used to identify a compound which interferes with the binding of MDM2 and p53. Since MDM2 is overexpressed in tumour cells and since binding of MDM2 to p53 appears to allow tumour cells to escape from p53-regulated growth, compounds that inhibit such binding would be useful as anti-cancer agents.

Sequence 64 AA;

0; Gaps Ouery Match 94.3%; Score 99; DB 19; Length 64; Best Local Similarity 100.0%; Pred. No. 2e-08; Matches 18; Conservative 0; Mismatches 0; Indels

1 PPLSQETFSDLWKLLPEN 18 ò

12 PPLSQETFSDLWKLLPEN 29 g Search completed: February 12, 2003, 10:34:44 Job time : 37 secs

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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

(without alignments)
121.770 Million cell updates/sec February 12, 2003, 10:33:46; Search time 15 Seconds Run on:

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19 Perfect score:

Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

283224 seqs, 96134422 residues Searched:

Total number of hits satisfying chosen parameters:

283224

length: 0 length: 2000000000 Minimum DB seq Maximum DB seq

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR_73:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		عدد			SUMMARIES	
Result No.	Score	Query	Length	80	QI	ption
	66	94.3	391	7	JC6193	tumor suppressor p
~	66	94.3	393	٦	DNHU53	cellular tumor ant
e	66	94.3	393	Н	806594	tumor
4	94	89.5	386	Н	S51648	cellular tumor ant
ស	98	81.9	393	~	JC6176	tumor suppressor p
ø	98	81.9	396	Н	JH0633	cellular tumor ant
7	81	77.1	363	٦	A29376	
80	74	70.5	381	N	538824	
6	74	70.5	390	Н	DNMS53	cellular tumor ant
10	72	9.89	391	Н	802192	cellular tumor ant
11	9		396	٦	JH0631	ımor
12	51.5	49.0	1009	~	S28857	qlutamate receptor
13	-	49.0	1009	7	JH0266	qlutamate receptor
14	20	47.6	230	7	C64396	precorrin-2 methyl
15	20	47.6	367	~	G86715	racemase [imported
16	20		447	N	E96672	Similar to Flavono
17	48	45.7	119	7	C49921	major intracellula
18	48	45.7	308	7	AD3262	methyltransferase
19	48		1004	N	B25039	outer cell wall pr
20	47	•	157	Н	VCTMPV	-
21	47	4	469	~	G84779	
22	47	₹.	487	~	D83027	hypothetical prote
23	47	₹.	489	~	T41241	H
24	47	44.8	926	7	G70327	isoleucine-tRNA li
25	46.5	44.3	250	~	T01604	hypothetical prote
56	46.5		485	~	C75460	
27	46	43.8	274	7	S74792	
28	46	43.8	293	Н	69	N
29	45.5	43.3	250	7	T49221	hypothetical prote

hypothetical prote	hypothetical prote	DNA-Binding Protei	conserved hypothet	conserved hypothet	L3 protein - bovin	hypothetical prote	hypothetical prote	NEF protein - simi	protein -	NEF protein - simi	protein -	NEF protein - simi	NEF protein - simi	protein -	1
E96796	AD2194	157557	B95064	D97931	P3WLB4	S03072	A82749	S61204	ASLJMS	S61205	S54852	S61206	S61208	S54853	S54849
7	~	7	7	~	Н	~	7	7	Н	7	7	7	7	7	~
367	405	837	211	211	113	114	134	209	211	238	239	239	244	246	250
42.9	42.9	42.9	42.4	42.4	41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9	41.9
45	45	45	44.5	44.5	44	44	44	44	44	44	44	44	44	44	44
30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45

ALIGNMENTS

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Cispecies: Oryctolagus cuniculus (domestic rabbit)
Cispecies: Nay, P.; Ronco, P.; de Fromentel, C.C.
Gene 185, 169-173, 1997
A;Title: CDNA cloning and immunological characterization of rabbit p53.
A;Rolecule type: mRNA
A;Reference number: JC6193; MUID:97208869; PMID:9055811
A;Accession: JC6193
A;Molecule type: mRNA
A;Residues: 1-391 <LEA>
A;Cross-references: EMBL:X90592; NID:g1532043; PIDN:CAA62216.1; PID:g1532044
C;Genetics: p53
C;Superfamily: cellular tumor antigen p53
C;Keywords: tumor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ..
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RESULT 1
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UNAULAR TUMOR antigen p53 [validated] - human
NyAlternate names: cellular phosphoprotein p53; oncoprotein p53; transformation suppress Cilbates: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 05-0ct-1988 #sequence revision 18-Nov-1994 #text change 15-Sep-2000
C;Accession: A2524; A43073; JT0436; S40773; S42669; A22837; A55060; A25397; S428905; ISB364; IYB850; IS2681; S60153
R;Lamb, P.; Crawford, L.
Mol. Cell. Biol. 6, 1379-1385, 1986
A;Title: Characterization of the human p53 gene.
A;Reference number: A25224; MUID:87064416; PMID:2946935
A;Residues: 1-33 < clam>A;Residues: 1-33 < clam>A;Reference number: JT0436; MUID:89108008; PMID:2905688
A;Title: A variation in the structure of the protein-coding region of the human p53 gene A;Reference number: JT0436; MUID:89108008; PMID:2905688
A;Residues: 1-333 < clam>A;Residues: 1-333 < clam>A;Residues: EMBL:M22898; NID:g189474

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A;Cross references: EMBL:X54156; NID:935213; PIDN:CAA38095.1; PID:935214
R;Yamada, Y.; Yoshida, T.; Hayashi, K.; Sekiya, T.; Yokota, J.; Hirohashi, S.; Nakatani,
Cancer Res. 51, 5800-5805, 1991
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A,Reference number: A44905, MUID:92034678, PMID:1933850
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Ajstaus: translated from GB/EMBL/DDBJ
Ajatus: translated from GB/EMBL/DDBJ
Ajatus: translated from GB/EMBL/DDBJ
Ajatus: translated from GB/EMBL/GBBJ
Ajatus: An Alu polymorphism intragenic to the TPS3 gene.
Ajatus: An Alu polymorphism intragenic to the TPS3 gene.
Ajatus: translated from GB/EMBL/GBJ
Ajatus: translated from GB/EMB
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A;Cross-references: EMBL:X60013; NID:g506438; PIDN:CAA42628.1; PID:g506439
A;Accession: 138086
A;Status: translated from GB/EMBL/DDBJ
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A;Cross-references: EMBL:X60014; NID:g506440; PIDN:CAA42629.1; PID:g506441
A;Accession: 138087
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A;Residues: 1-247,'Q',249-393 <F06>
A;Cross-references: EMBL:X60015; NID:9506442; PIDN:CAA42630.1; PID:9506443
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A.Cross-references: EMBL:X60016; NID:9506444; PIDN:CAA42631.1; PID:9506445
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A;Molecule type: mRNA
A;Molecules: 1-247, (2), 249-393 <PG8>
A;Cross-references: EMBL:X60017; NID:g506446; PIDN:CAA42632.1; PID:g506447
                                                                                                                                                                                                                    A;Molecule type: mRNA
A;Residues: 1-192,'R',194-393 <F02>
A;Cross-references: EMBD:X60011; NID:9506434; PIDN:CAA42626.1; PID:9506435
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     A; Cross-references: EMBL:X60010; NID:g506432; PIDN:CAA42625.1; PID:g506433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Cross-references: EMBL:X60012; NID:g506436; PIDN:CAA42627.1; PID:g506437
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A;Cross-references: EMBL:X60019; NID:g506450; PIDN:CAA42634.1; PID:g506451
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A;Residues: 246-247, 'W', 249-250 <YAM>
A;Residues: 246-247, 'W', 249-250 <YAM>
A;Cross-references: GB:S63157; NID:9237829; PIDN:AAB20140.1; PID:9237830
A;Note: sequence extracted from NCBI backbone (NCBIN:63157, NCBIP:63158)
A;Note: mutation from a liver metastasis of a gastric cancer
R;Hensel, C.H.; Xiang, R.H.; Sakaguchi, A.Y.; Naylor, S.L.
                                                               A, Note: deletion of a C nucleotide causes a frameshift at position 566
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A;Residues: 1-71,'P',73-162,'H',164-393 <F09>
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A;Molecule type: mRNA
A;Residues: 1-393 <F03>
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A,Status: translated from GB/EMBL/DDBJ
A,Molecule type: mRNA
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A; Status: translated from GB/EMBL/DDBJ
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A, Molecule type: DNA
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A,Status: translated from GB/EMBL/DDBJ
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A; Residues: 1-393 <CHU>
A; Cross-references: EMBL:X54156; NID:g35213; PIDN:CAA38095.1; PID:g35214
A; Cross-references: EMBL:X54156; NID:g35213; PIDN:CAA38095.1; PID:g35214
B; Matlashewski, G.; Lamb, P.; Pim, D.; Peacock, J.; Crawford, L.; Benchimol, S.
EMBO J. 3, 3257-3262, 1984
A; Title: Isolation and characterization of a human p53 cDNA clone: expression of the hum A; Reference number: 842669; MUID:85126934; PMID:6396087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A,Molecule type: mRNA
A;Residues: 1-71, 'P. 73-393 <ZAK>
A;Residues: 1-71, 'P. 73-393 <ZAK>
A;Cross-references: EMBL:X02469; EMBL:M60950; NID:g35209; PIDN:CAA26306.1; PID:g35210
R;Harlow, E., Williamson, N.M.; Ralston, R.; Helfman, D.M.; Adams, T.E.
Mol. Cell. Biol. 5, 1601-1610, 1985
A;Title: Molecular cloning and in vitro expression of a cDNA clone for human cellular tu
A;Reference number: A55060; MUID:85267676; PMID:3894933
A;Note: this 72-Arg allele appears to be about 5 times more frequent than the 72-Pro all
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A;Molecule type: mRNA
A;Residues: 1-189,'LLSILSEWKEICVWSIMMTETLFDIVWWCPMSRLRLALT','VPPSTTTTCVTVPAWAA' <F01>
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A; Cross-references: EMBL:X01405; NID:g35215; PIDN:CAA25652.1; PID:g642241
A; Cross-references: EMBL:X01405; NID:g35215; PIDN:CAA25652.1; PID:g642241
B; Zakut-Houri, R.; Bienz-Tadmor, B.; Givol, D.; Oren, M.
EMBO J. 4, 1251-1255, 1985
A; Title: Human p53 cellular tumor antigen: cDNA sequence and expression in COS cells.
A; Reference number: A22837; MUID:85230577; PMID:4006916
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A; Residues: 1-71, PP. 773-272, 'H', 274-393 < HAR>
A; Cross-references: GB: K03199; NID: g189479; PIDN: AAA59989.1; PID: g189479
A; Experimental source: clone pR4-2, cell line A431
R; Harris, N.; Brill, E.; Shohat, O.; Prokocimer, M.; Wolf, D.; Arai, N.; Rotter, V. Mol. cell. Biol. 6, 4650-4656; 1986
A; Fitle: Molecular basis for heterogeneity of the human p53 protein.
A; Reference number: A93086; MUID: 87089926; PMID: 3025664
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A,Residues: 1-71, 'P',73-78,'T',80-393 «HAR2>
A,Residues: 1-71,'P',73-78,'T',80-393 «HAR2>
A,Residues: 1-71,'P',73-78,'T',80-393 «HAR2>
A,Gross-references: EMBL:M14695; NID:g339816
A,Experimental source: clone p53-H-19, transformed hybridoma 8V-80 cell line
R,Matlashewski, G.J.; Tuck, S.; Pim, D.; Lamb, P.; Schneider, J.; Crawford, L.V.
Mol. Cell. Biol. 7, 961-963, 1987
A,Title: Primary structure polymorphism at amino acid residue 72 of human p53.
A,Reference number: 842452,
A,Accession: 842452.
                                                                                                                                                                                                               A,Cross-references: EMBL:M22898; NID:g189474; PIDN:AAAS9988.1; PID:g189476
A,Note: this 72-Pro allele was found in both normal and malignant cell lines
R;Chumakov, P.M.; Amazov, V.P.; Jenkins, J.R.
submitted to the EMBL Data Library, August 1990
A;Reference number: $40773
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A;Cross-references: EMBL:M14694; NID:9339813; PIDN:AAAG1211.1; PID:9339814
A;Experimental source: clone p53-H-1, transformed hybridoma SV-80 cell line
A;Accession: B25397
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A;Residues: 66-71, 'P',73-79 <MKI2>
A;Experimental source: clone lambda C113
A;Note: 72-Cys was also found, and appears to represent a polymorphism
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A;Residues: 66-79 <MKI3>
A;Experimental source: clone J6K
Experimental; P.U.; Allan, G.U.; Shanahan, F.; Vousden, K.H.; Crook, T.
EMBO J. 10, 2879-2887, 1991
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A,Reference number: 138082; MUID:92007731; PMID:1915267
                                                                                                                   A, Molecule type: DNA
A, Residues: 1-71,'P',73-393 <BUC2>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Accession: S42453
                                                                     A; Accession: JT0436
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cellular tumor antigen p53 - bovine
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nes 16; Conserva
                                                                                                                                                                                                                                                                                                        A Status: preliminary
A Molecule type: mRNA
A;Residues: 1-386 <DEQ>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Accession: JC6176
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A, Residues: 1-393 <RIG>
A, Residues: 1-394 <RIG>
A, Residues: 1-395 <RIG>
A, Crose-references: EMBLS.X16384, NID:922795; PIDN:CAA34420.1; PID:922796
C, Superfamily: cellular tumor antigen p53
C, Reywords: apoptosis; cell division control; DNA binding; homotetramer; nucleus; phosph P;176,179,238,247/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;392/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
                                                                                                                                                                                                                                                                                                     A; Molecule type: DNA
A; Residues: 274-277, S',279-282 < HEN2>
A; Residues: 274-277, S',279-282 < HEN2>
A; Cross-references: GB $41977, NID:91679932; PIDN:AAB19325.1; PID:9232816
R; Chow, V. T.; Quek, H.H.; Tock, E.P.C.
Cancer Lett. 73, 141-148, 1993
A; Title: Alternative splicing of the p53 tumor suppressor gene in the Molt-4 T-lymphobla
A; Reference number: IS2681; MUID:94036762; PMID:8221626
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Molecule type: mRNA
A; Residues: 327-311, DQTSFQKENC' <CHO>
A; Cross-references: GB:S66666; NID:9436292; PIDN:AAB28601.1; PID:9436293
A; Note: mutant sequence with altered splicing and termination expressed in Molt-4 T-lymp
A; Note: mutant sequence with altered splicing and termination expressed in Molt-4 T-lymp
A; Note: mutant sequence with altered splicing and termination expressed in Molt-4 T-lymp
Mol. Gen. Genet. 249, 425-431, 1995
A; Title: Mapping of linear epitopes recognized by monoclonal antibodies with gene-fragme
A; Reference number: S60151; MUID:96133682; PMID:8552047
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: DNA
A;Residues: 3-44 <PET>
R;Dang, C.V.; Lee, W.M.F.
J. Blol. Chem. 264, 18019-18023, 1989
A;Title: Nuclear and nucleolar targeting sequences of c-erb-A, C-myb, N-myc, p53, HSP70,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 cellular tumor antigen p53 - green monkey
C;Species: Cercopithecus aethiops (green monkey, grivet)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: 806594
R;Rigaudy, P.; Eckhart, W.
Nucleic Acids Res. 17, 8375, 1989
A;Title: Nucleotide sequence of a cDNA encoding the monkey cellular phosphoprotein p53.
                          A,Title: Use of the single strand conformation polymorphism technique and PCR to detect A,Reference number: 158354; MUID:91296386; PMID:1648702
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                                                                                                                                               A; Modecule type: DNA
A; Residues: 244-247, 'W', 249-252 <HEN1>
A; Cross-references: GB: S41969; NID: g1679931; PIDN: AAB19324.1; PID: g232814
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94.3%; Score 99; DB 1; Length 393; 100.0%; Pred. No. 3e-07; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                       A,Accession: 178850
A,Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                     A; Status: preliminary; translated from GB/EMBL/DDBJ
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Best Local Similarity 100.
Matches 18; Conservative
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   Oncogene 5, 1067-1071, 1991
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Matches 18; Conserv
                                                                                           Accession: 158354
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S51648

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A; Modecule type: mRNA
A; Residues: 1-393 <LEE>
A; Residues: 1-393 <LEE>
A; Cross-references: GB:U50395; NID:g1842229; PIDN:AAC53040.1; PID:g1842230
C; Comment: This protein is a multimer, it plays the central role in a complex DNA damage iption, and recombination by protein/protein interactions.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A;Cross-references: EMBL:X81704; NID:g602332; PIDN:CAA57348.1; PID:g602333
C;Superfamily: cellular tumor antigen p53
C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; phosphoprotein; F;168,171,231,235;Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;385/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
NiAlternate names: tumor-suppressor protein p53
C;Species: Bos primigenius taurus (cattle)
C;Species: Bos primigenius taurus (cattle)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: S51648
R;Dequiedt, F.; Willems, L.; Burny, A.; Kettmann, R.
submitted to the EMBL Data Library, September 1994
A;Description: Nucleotide sequence of the ovine p53 tumor-suppressor gene cDNA and its A;Reference number: S51648
A;Accession: S51648
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NyAlternate names: tumor-suppressor protein p53
Cispecies: Mesocricetus auratus (golden hamster)
Cispecies: Mesocricetus auratus (golden hamster)
Cispecies: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
Cisaccesion: UH6633
Rilegros, Y.; McIntyre, P.; Soussi, T.
Gene 112, 247-250, 1992
A; At-250, 199
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A;Reference number: JC6176; MUID:97183659; PMID:9031625
A;Contents: liver
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 393;
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89.5%; Score 94; DB 1; Length 386;
Best Local Similarity 94.4%; Pred. No. 1.7e-06;
Matches 17; Conservative 0; Mismatches 1; Indels
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Pred. No. 2.8e-05;
0; Mismatches 1;
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C,Superfamily: cellular tumor antigen p53
C,Keywords: liver; tumor
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A;Residues: 1-396 <LEG>
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A; Molecule type: mRNA
A; Residues: 1-381 cARA>
A; Cresidues: 1-381 cARA>
A; Cresidues: 1-381 cARA>
A; Cresidues: 1-381 cARA>
A; Cresidues: 1-381 cARA>
B; Han, K.A.; Kulesz-Martin, M.F.
Nucleic Acids Res. 20, 1979-1981, 1992
Nucleic Acids Res. 20, 1979-1981, 1992
A; Title: Alternatively spliced p53 RNA in transformed and normal cells of diff-rent tiss
A; Reference number: 835478; MUID:92253421; PMID:1579500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Cross-references: EMBL:M13874; NID:g200202; PIDN:AAA39883.1; PID:g200203
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, July 1388
C;Comment: This sequence, produced by alternative splicing of the tenth intron, lacks th
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A;Cross-references: GB:X00876; NID:g871420; PIDN:CAA25420.1; PID:g871421; GB:X01237; GB
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Filt0-192/Region: conserved region II
Filt0-192/Region: conserved region III
Filt0-192/Region: conserved region IV
Filt0-192/Region: conserved region IV
Filt0-193-248/Region: conserved region IV
Filt0-193/Region: conserved region V
Filt0-193/Region: tetramer association
Filt0-193/Region: tetr
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(c)Species: Mus musculus (house mouse)
(c)Species: Mus musculus (house mouse)
(c)Date: 28-Aug-1985 #sequence_revision 04-Oct-1996 #text_change 11-May-2000
(c)Accession: A22739, 806336, A02684, 838822, 838823, 840014; I48703
(R)Bienz, B.; Zakut-Houri, R.; Givol, D.; Oren, M. M. Shenz, B.; Zakut-Houri, R.; Givol, D.; Oren, M. A. Xille: Analysis of the gene coding for the murine cellular tumour antigen p53.
A)Reference number: A22739; MUID:85027173; PMID:6092064
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A;Title: Primary structure of DNA complementary to murine oncoprotein p53 mRNA.
A;Reference number: S06336; MUID:88221682; PMID:3329909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A,Molecule type: mRNA
A,Residues: 1-134, V',136-390 <CHU>
R;Zakut-Houri, R.; Oren, M.; Bienz, B.; Lavie, V.; Hazum, S.; Givol, D.
Nature 306, 594-597, 1983
A;Title: A single gene and a pseudogene for the cellular tumour antigen p53.
A;Reference number: A02684; MUID:84068204; PMID:6646235
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                                                                                                                                                                                                                                                                                                                                                                                                                                      A; Status: nucleic acid sequence not shown; translation not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           s not known.
C;Superfamily: cellular tumor antigen p53
C;Keywords: alternative splicing; phosphoprotein; zinc
E;1-44/Domain: transcription activation #status predicted <TRA>
E;16-26/Region: conserved region I
E;99-289/Domain: DNA-binding core #status predicted <DBC>
F;108-121/Region: L loop
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A;Residues: 1-159,'H',161-167,'G',169-233,'I',235-390 <ZAK>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Status: not compared with conceptual translation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             cellular tumor antigen p53 - mouse
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A; Residues: 1-381 <HAN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
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A; Accession: S38824
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Best Local 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   cellular tumor antigen p53 - African clawed frog
C;Species: Xenopus laevis (African clawed frog)
C;Date: 10-Sep-1999 #sequence revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession A29376; S61531; $72313; 151639
R;Soussi, T.; de Fromentel, C.C.; Mechali, M.; May, P.; Kress, M.
Oncogene 1, 71-78, 1987
A;Title: Cloning and characterization of a CDNA from Xenopus laevis coding for a protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Molecule type: mRNA
A; Residues: 1-363 <SOUS
A; Residues: 1-363 <SOUS
A; Cross-references: EMBL: X05191; NID: 964961; PIDN: CAA28821.1; PID: 964962
A; Cross-references: EMBL: X05191; NID: 964961; PIDN: CAA28821.1; PID: 964962
B; Hoever, M.; Clement, J.H.; Wedlich, D.; Montenarh, M.; Knoechel, W.
Oncogene 9, 109-120, 1994
Oncogene 9, 109-120, 1994
A; Title: Overexpression of wild-type p53 interferes with normal development in Xenopus 1
A; Reference number: 151639; MUID: 94134403; PMID: 8302570
                                                                                                                                                                                C;Superfamily: cellular tumor antigen p53
C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nucleus; phosph
F;179,182,241,245/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;395/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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C;Keywords: apoptosis; cell division control; DNA binding; homotetramer; nucleus; phosph P;150,153,213,217/Binding site: zinc (Cys, His, Cys, Cys) #status predicted P;362/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
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Cispecies: Mus musculus (house mouse)
Ciplate: 13-Jan-1995 #sequence_revision 13-Jan-1995 #text_change 23-Jul-1999
CiAccession: 838824; 835478
Mol. A. Nomura, D.; Yokota, K.; Wolf, D.; Brill, E.; Shohat, O.; Rotter, V. Mol. Cell. Biol. 6, 3232-3239, 1986
A; Title: Immunologically distinct p53 molecules generated by alternative splicing. A; Reference number: $38822; MuID:8706440; PMID:3023970
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A;Residues: 1-293,295-363 <HOE>
A;Residues: 1-293,295-363 <HOE>
A;Cross-references: EMBL:X77546; NID:g468513; PIDN:CAA54672.1; PID:g468514
R;Hoever, M.; Clement, J.; Wedlich, D.; Montenarh, M.; Knochel, W.
submitted to the EMBL Data Library, March 1994
A;Reference number: S72313
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A,Residues: 1-51,'S',53-70,72-293,295-363 <HOW>
A,Cross-references: EMBL:X77546; NID:g468513; PIDN:CAA54672.1; PID:g468514
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     A; Cross-references: GB: M75144; NID: g191414; PIDN: AAA37085.1; PID: g191415
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                                             ;Experimental source: kidney, strain MP1
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Matches 14; Conservative
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                                                                                                                                            A;Gene: p53
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A;Cross-references: EMBL:X13058; NID:956828; PIDN:CAA31457.1; PID:956829
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A; Cross-references: EMBL: Z17238
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A;Molecule type: mRNA
A;Residues: 1-47, 'R', 49-78 'QW', 82-390 <RES>
A;Cross-references: EMBL:X00741, NID:953570; PIDN:CAA25323.1; PID:953571
C;Comment: This DNA-binding protein plays an essential role in the regulation of cell di
C;Comment: The tetramer association region may exhibit a beta-turn, beta-sheet, beta-tur
C;Superfamily: cellular tumor antigen p53
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F;1-44/pomain: transcription activation #status predicted <TRA>
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A; Cross-references: EMBL:M13873; NID:g200200; PIDN:AAA39882.1; PID:g200201
A; Cross-references: EMBL:M3873; NID:g200200; PIDN:AAA39882.1; PID:g200201
B; Jenkins, J.R.; Rudge, K.; Redmond, S.; Wade-Evans, A.
Nucleic Acids Res. 12, 5609-5626, 1984
A; Title: Cloning and expression analysis of full length mouse cDNA sequences encoding A; Reference number: 148703; MUID:84272240; PMID:6379601
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F;173,176,235,239/Binding site: zinc (Cys, His, Cys, Cys) #status predicted
F;1312/Binding site: phosphate (Ser) (covalent) (by cdc2 kinase) #status predicted
F;389/Binding site: phosphory1-RNA (Ser) (covalent) #status predicted
          A;Cross-xeferences: GB:X01237; GB:K01700; NID:g53575
R;Arai, N.; Nomura, D.; Yokota, K.; Wolf, D.; Brill, E.; Shohat, O.; Rotter, V.
Mol. Cell. Biol. 6, 3323-3239, 1986
A;Title: Immunologically distinct p53 molecules generated by alternative splicing. A;Reference number: S38822; MUID:87064640; PMID:3023970
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Nucleic Acids Res. 16, 11384, 1988
A;Title: Nucleotide sequence of a CDNA encoding the rat p53 nuclear oncoprotein.
A;Reference number: S02192; MUID:89083585; PMID:3060862
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A; Residues: 1-167, 'G', 169-233,'I', 235-390 <ARA2>
A; Crosel-references: BMBL:MI333
R; Arai, N.; Nomura, D.; Yokota, K.; Wolf, D.; Brill, E.; Shohat, O.; Rotter, V. submitted to the EMBL Data Library, July 1988
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N'Alternate names: gene p53 protein; nuclear oncoprotein p53
C;Species: Rattus norvegicus (Norway rat)
C;Date: 10-Sep-1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
C;Accession: S02192; S41149
                                                                                                                                                                                                                A;Molecule type: mRNA
A;Residues: 1-390 <ARA1>
A;Cross-references: EMBL:M13872; NID:g200198; PIDN:AAA39881.1; PID:g200199
A;Accession: S38823
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F;99-289/Domain: DNA-binding core #status predicted <DBC>
F;108-121/Region: L1 loop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          F;233-248/Region: L3 loop
F;267-283/Region: conserved region V
F;313-319/Region: nuclear location signal
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F;168-178/Region: conserved region III
F;231-252/Region: conserved region IV
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A, Molecule type: mRNA
A, Residues: 1-391 <SOU>
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Glutamate receptor delta-1 chain precursor - rat
NyAlternate names: Asinate receptor
NyAlternate names: Asinate receptor
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 07-Apr-1994 #sequence_revision 07-Apr-1994 #text_change 20-Aug-1999
C;Accession: S28857; S31222
C;Accession: S28857; K.; Laurie, D.J.; Koehr, G.; Herb, A.; Seeburg, P.H.; Wisden, W.FEBS Lett. 315, 318-322, 1993
A;Pitle: The rat delta-1 and delta-2 subunits extend the excitatory amino acid receptor
A;Reference number: S28857; MUID:93138096; PMID:8422924
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Introns: 25/2; 32/3; 123/3; 185/1; 259/2; 305/1; 329/3; 365/2 C; Superfamily: cellular tumor antigen p53 C; Superfamily: cellular tumor antigen p53 C; Superfamily: cell division control; DNA binding; homotetramer; nucleus; phosph F;174,177,236,240/Binding site: zinc (Cys, His, Cys, Cys) #status predicted F;390/Binding site: phosphoryl-RNA (Ser) (covalent) #status predicted
                                                                                                                                                                                    A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Status: preliminary; nucleic acid sequence not shown; translation not shown A;Molecule type: DNA A;Residues: 1-173, W, 175-391 (HUL) A;Cross-references: EMBL:L07909 A;Note: the nucleotide sequence was submitted to the EMBL Data Library, December 1992 C;Genetics:
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A;Title: Rainbow trout p53: cDNA cloning and biochemical characterization.
A;Reference number: JH0631; MJID:92210006; PMID:1339362
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              cellular tumor antigen p53 - rainbow trout
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Date: 10-Sep_1999 #sequence_revision 10-Sep-1999 #text_change 10-Sep-1999
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68.6%; Score 72; DB 1; Length 391;
Best Local Similarity 93.3%; Pred. No. 0.0037;
Matches 14; Conservative 0; Mismatches 1; Indels
R;Hulla, J.E.; Schneider, R.P.
Nuclèic Acids Res. 21, 713-717, 1993
A;Title: Structure of the rat p53 tumor suppressor gene.
A;Reference number: S41149; MUID:93181268; PMID:8441680
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Best Local Similarity
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C;Accession: JH0266
C;Accession: JH0266
C;Accession: JH0266
Biochem. Biophys. Res. Commun. 183, 886-892, 1992
A;Title: Molecular cloning of a cDNA encoding a novel member of the mouse glutamate recent and a common of a cDNA encoding a novel member of the mouse glutamate recent are number. JH0266; MUID: 92198486; PMID: 1372507
                                                                                                                                                                                                                                                                                             C; Superfamily: glutamate receptor; glutamate, receptor homology
C; Keywords: glycoprotein; ion channel; neurotransmitter receptor; transmembrane protein
C; Keywords: glycoprotein; ion channel; neurotransmitter receptor; transmembrane protein
F;1-15/Domain: signal sequence #status predicted <SIG>
F;16-1009/Product: glutamate receptor delta-1 chain #status predicted <MAT>
F;55-584/Domain: transmembrane #status predicted <TM1>
F;601-621/Domain: transmembrane #status predicted <TM3>
F;32-654/Domain: transmembrane #status predicted <TM3>
F;33-654/Domain: transmembrane #status predicted <TM3>
F;33-654/Domain: transmembrane #status predicted <TM3>
F;33-654/Domain: transmembrane #status predicted <TM3>
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A,Residues: 11.009 
A,Residues: 11.009 
A,Residues: 11.009 
A,Rorasare references: DBD1:D10171; NID:g220417; PIDN:BAA01041.1; PID:g220418
C,Comment: Glutamate receptor channels mediate most of the fast excitatory synaptic tran
C,Superfamily: glutamate receptor; glutamate receptor homology
C,Teywords: glytoprotein; neurotransmitter receptor; phosphoprotein; transmembrane prote
F;16-1009/Product: glutamate receptor channel delta-1 chain #status predicted <GLU>
F;16-1009/Product: glutamate receptor channel delta-1 chain #status predicted <GLU>
F;16-1009/Product: glutamate receptor homology <GRH>
F;10-1009/Product: glutamate
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C; Species: Methanococcus jannaschii
C; Date: 13. Sep-1996 #sequence_revision 13. Sep-1996 #text_change 21-Jul-2000
C; Accession: C64396
R; Bult, C.J.; White, O.; Olsen, G.J.; Zhou, L.; Fleischmann, R.D.; Sutton, G.G.; Blake,
? Reich, C.I.; Overbeek, R.; Kirkness, E.F.; Weinstock, K.G.; Merrick, J.M.; Glodek, A.;
rson, J.D.; Śadow, P.W.; Hanna, M.C.; Cotton, M.D.; Roberts, K.M.; Hurst, M.A.
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C:Species: Mus musculus (house mouse)
C:Date: 30-Jun-1992 #sequence_revision 30-Jun-1992 #text_change 20-Jun-2000
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                                                                                                                                                                       A;Molecule type: mRNA
A;Residues: 83-1009 <SPR>
A;Cross-references: EMBL:Z17238; NID:956285; PIDN:CAA78936.1; PID:956286
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R,Sprengel, R.
submitted to the EMBL Data Library, October 1992
A,Reference number: S31222
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        47.48;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 9; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Best Local Similarity
2, Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity
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                                                                                                                                   A; Accession: S31222
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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Science 273, 1058-1073, 1996
A;Authors: Kaine, B.P.; Borodovsky, M.; Klenk, H.P.; Fraser, C.M.; Smith, H.O.: Woese, C.A;Title: Complete genome sequence of the methanogenic archaeon, Methanococcus jannaschii A;Reference number: A64300; MUID:96337999; PMID:8688087
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  C,Accession: G86715
R;Bolotin, A.; Wincker, P.; Mauger, S.; Jaillon, O.; Malarme, K.; Weissenbach, J.; Ehrli
Genome Res. 11, 731-753, 2001
A;Title: The complete genome sequence of the lactic acid bacterium Lactococcus lactis ss
A;Reference number: A86625; WUID:21235186; PMID:11337471
                                                                                                                                                                                                                 A;Residues: 1-230 <BUL>
A;Cross-references: GB:U67522; GB:L77117; NID:g2826315; PIDN:AAB98764.1; PID:g.592306;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Residues: 1.367 <STO>
A;Cross-references: GB:AE005176; PID:g12723640; PIDN:AAK04825.1; GSPDB:GN00146
A;Experimental source: strain IL1403
                                                                                                                         A,Accession: C64396
A,Status: preliminary; nucleic acid sequence not shown; translation not shown
A,Molecule type: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Lactococcus lactis subsp. lactis
C;Date: 23-Mar_2001 #sequence_revision 23-Mar-2001 #text_change 03-Aug-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             racemase [imported] - Lactococcus lactis subsp. lactis (strain IL1403)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ·.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            47.6%; Score 50; DB 2; Length 367;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                      47.6%; Score 50; DB 2;
55.6%; Pred. No. 4.5;
tive 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 7.5;
3; Mismatches
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                                                                                                                                                                                                                                                                                                                                       A;Start codon: GTG
C;Superfamily: precorrin-3 methylase
                                                                                                                                                                                                                                                                                                             position: FOR691742-692434
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      101 PTLYSTFSYVWKLLKERG 118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         52.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2 PLSQETFSDLWKLLPENG 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            54 PFYTSETFADSWKILEE 70
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GenCore version 5.1.3
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OM protein - protein search, using sw model

Run on:

February 12, 2003, 10:32:56 ; Search time 11 Seconds (without alignments) 71.641 Million cell updates/sec

US-09-403-440A-2 105 1 PPLSQETFSDLWKLLPENG 19 Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

112892 segs, 41476328 residues Searched:

112892 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

SwissProt_40:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		de			SUMMARIES	
Result		Query				
No.	Score	Match	Length	DB	QI	Description
-	66	94.3	386	Н	P53 PIG	Q9tub2 sus scrofa
7	66	94.3	391	-	P53 CAVPO	O9wur6 cavia porce
m	66	94.3	391	Н		-
4	66	94.3	393	-	P53_CERAE	_
S	66	94.3	393	Н		
9	66	94.3	393	Н	ìα	macac
7	66		393	Н	iń	тасаса
80	66		393	Н	23	Q9ttal tupaia glis
6	94		382	-	P53_SHEEP	ovis an
10	94	89.5	386	Н	P53_BOVIN	Q29628 bos taurus
11	94		391	Н	- 1	marmot
12	90		381	-	23	canis
13	90	85.7	386	Н		P41685 felis silve
14	98		393	-	P53_CRIGR	cricet
15	98	81.9	396	П	P53_MESAU	
16	81		363	Н	P53_XENLA	P07193 xenopus lae
17	74	70.5	390	7	P53_MOUSE	
18	72		391	Н	P53_RAT	P10361 rattus norv
19	9		396	н	P53_ONCMY	P25035 oncorhynchu
50	'n,		367	Н		
21	53.5	급.	4	-	P53_XIPHE	
22	m	-	342	-1	P53_XIPMA	
23	20	47.6	m	-		
24	48	ď.	119	Н	IPI_BACSU	
25	48		1004	Н	SLPO_BACBR	bacillu
56	47	4	156	-	COAT_PMMV2	pepper
27	47	4.	156	-	COAT_PMMVI	pepper
28	47	4.	156	-	COAT_PMMVS	
23	47	4.	156	-	COAT_TOMKO	tomato
30	47		158	٦	COAT_PMMV0	
31	4.1	4	~	-	- 1	
32	47	44.8	926	н	۷,	
33	46	•	0	н	YA24_SYNY3	

P52633 mus musculu Q13188 homo sapien	P08343 bovine papi P58879 methanopyru	P05861 simian immu P31818 simian immu	P11262 simian immu	P10360 gallus gall	Q00976 kluyveromyc	P39847 bacillus su
STA6_MOUSE STK3_HUMAN	VL3 BPV4 HIS7 METKA	NEF SIVMK	NEF_SIVML	P53 CHICK	CC68 KLULA	PPS3_BACSU
						-
837	113	207	263	367	1033	2555
42.9	41.9	41.9	41.9	41.9	41.9	41.9
45 44.5	4 4 4 4	4 4 4 4	4 4 4 4	4 4	44	44
3.4 3.5	36	3 8 8 6 8 6	40	4 4 2	4 4	45

ALIGNMENTS

DR Fram; PrOUBYO; PS3; 1. DR PRINTS; PRO0386; PS3SUPPRESSR. DR ProDom; PD002681; PS3; 2.

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                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BCl-2
                BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- SUBUNIT: Binds DNA as a homotetramer (By similarity).
-!- SUBCELUIAR LOCATION: Nuclear.
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               D'erchia A.M., Pesole G., Tullo A., Saccone C., Sbisa E.; "Guinea pig p53 mRNA: identification of new elements in coding and
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cavia porcellus (Guinea pig).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Rodentia, Hystricognathi, Caviidae, Cavia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nučlear protein; Phosphorylation; Apoptosis.

1 44 TRANSCRIPTION ACTIVATION (ACIDIC).

DNA BIND 100 290 BY SIMILARITY.
                                                                                                                                                    ;
                                                                                                              94.3%; Score 99; DB 1; Length 386;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   untranslated regions and their functional and evolutionary
                                                                                                                                                    0; Indels
318 349 OLIGOMERIZATION.
361 380 BASIC (REPRESSION OF DNA-PACHER 1304 316 NUCLEAR LOCALIZATION SIGNA PHOSPHORYLATION (BY SIMILA 386 AA; 42862 MW, A4C3D88REBDF55162 CRC64;
                                                                                                                                                                                                                                                                                                                                                            16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
15-UTN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                7.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OLIGOMERIZATION.
                                                                                                                                                                                                                                                                                                                          391 AA.
                                                                                                                            100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         IN MANY TYPES OF CANCER.
-!- SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                          PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=99265972; PubMed=10331945;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      HSSP, P04637; 1YCS.
InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
Probom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; AJ009673; CAB43196.1; -.
                                                                                                                                                                                        1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                          12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Genomics 58:50-64(1999).
                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                          STANDARD;
                                                                                                                              Local Similarity
nes 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=10141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                323
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   implications."
                                                                                                                                                                                                                                                                                                                        P53 CAVPO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DOMĀIN
                                                                          SEQUENCE
                                                                                                                Query Match
                                                        MOD RES
                  DOMAIN
                                      OMAIN
                                                                                                                                                      Matches
                                                                                                                                                                                                                                                                                                        P53_CAVPO
                                                                                                                                                                                                                                                                                      RESULT 2
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    84448
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NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE=97208869; PubMed=9055811; le Goas F., May P., Ronco P., Caron de Fromentel C.; "CDNA cloning and immunological characterization of rabbit p53.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 orgo.
Orgocolagus cuniculus (Rabbit).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Lagomorpha, Leporidae, Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HSSP; P04637; 1XCS.
InterPro; IRRO02117; PS3.
Pfam; PF00870; PS3; 1.
PRINTS; PR00386; PS3SUPPRESSR.
ProDom; P0002861; PS3; 1.
PROSITE; PS00348; PS3; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Nuclear protein; Phosphorylation; Apoptosis.

1 43 TRANSCRIPTION ACTIVATION (ACIDIC).
                                                                                                                                                                                           ..
0
                                                     PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                     DB 1; Length 391; 7.3e-08;
                                                                            321D40702383573E CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1997 (Rel. 35, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BY SIMILARITY.
OLIGOMERIZATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OF TRANSPORMED CELLS. P53 IS FREQUENTLY IN MANY TYPES OF CANCER. SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                        0; Mismatches
                                                                                                                                                                 Pred. No.
                                                                                                                                     94.3%; Score 99;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-NOV-1997 (Rel. 35, Created)
                                                                                  43287 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; X90592; CAA62216.1; -.
                                                                                                                                                                 100.08;
                                                                                                                                                                                                                                                 1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                           12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
                                                     390
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=New Zealand;
                                                                                                                                                         Local Similarity
hes 18; Conserv
                                                                               391 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             323
366
308
390
  366
309
390
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DOMAIN
                                                     MOD RES
SEQUENCE
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MOD RES
                                                                                                                                     Query Match
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                               DOMAIN
                                                                                                                                                                                           Matches
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393 AA; 43696 MW; 9ED285C9A7855D6E CRC64;

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SQ SEQUENCE
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                                                                             Matches
                                                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation-the European Bioinformatics Institute. There are no restrictions on its most by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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DOMAIN 1 44 TRANSCRIPTION ACTIVATION (ACIDIC).

DOMAIN 292 915 SIMILARITY.

DOMAIN 368 387 BASIC (REPRESSION OF DNA-BINDING).

DOMAIN 313 323 PHOSPHORYLATION (BY SIMILARITY).

PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        phosphoprocein p33.";
Nucleic Acids Res. 17:875-8375(1989).

**INUCLION: Acts as a tumor suppresor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BCI-2
                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               expression.
SUBDNIT: Binds DNA as a homotetramer (By similarity).
SUBCELLULAR LOCATION: Nuclear.
DISBASE: PS1 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. PS3 IS FREQUENTLY MUTATED OR INACTIVATED
IN MANY TYPES OF CANCER.
                                                                                                                                                                                                                                                                                                                                                                                               Cercopithecus aethiops (Green monkey) (Grivet).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Cercopithecidae,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIINE=90045967; Pubmed=2530498;
Rigaudy P., Eckhardt W.;
"Nucleotide sequence of a cDNA encoding the monkey cellular
                                     94.3%; Score 99; DB 1; Length 391; 100.0%; Pred. No. 7.3e-08; Live 0; Mismatches 0; Indels
391 AA; 43435 MW; 86BD5B8D0B726525 CRC64;
                                                                                                                                                                                                                                                                                                                                                            Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                 01-07N-1990 (Rel. 13, Created)
01-07NV-1990 (Rel. 13, Last sequence update)
15-07UN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                              393 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cercopithecinae, Cercopithecus.
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                                                                                                                      1 PPLSQETFSDLWKLLPEN 18
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InterPro; IPR002117; P53.
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PROSITE; PS00348; P53; 1.
                                                            Local Similarity 100.
Les 18; Conservative
                                                                                                                                                                                                                                                              STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Harlow E., Williamson N.M., Ralston R., Helfman D.M., Adams T.E.; "Molecular cloning and in vitro expression of a cDNA clone for human
                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE-85230577; PubMed-4006916;
Zakut-Houri R., Bienz-Tadmor B., Givol D., Oren M.;
"Human p53 cellular tumor antigen: cDNA sequence and expression in
COS cells.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cellular tumor antigen p53 (Tumor suppressor p53) (Phosphoprotein p53) (Antigen NY-CO-13).
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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MEDLINE=87089826; PubMed=3025664;
Harris N., Brill E., Shohat O., Prokocimer M., Wolf D., Arai N.,
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Farrell P.J., Allan G., Shanahan F., Vousden K.H., Crook T.;
"ps3 is frequently mutated in Burkitt's lymphoma cell lines.";
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Matlashewski G., Lamb P., Pim D., Peacock J., Crawford L.,
94.3%; Score 99; DB 1; Length 393; 100.0%; Pred. No. 7.3e-08; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                           P04637; Q16848; Q9UBI2;
13-AUG-1987 (Rel. 05, Created)
01-MAR.1989 (Rel. 10, Last sequence update)
15-UUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                   393 AA.
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MEDLINE=87064416; PubMed=2946935;
Lamb P., Crawford L.;
"Characterization of the human p53 gene.";
MOl. Cell. Biol. 6:1379-1385(1986).
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Mol. Cell. Biol. 5:1601-1610(1985).
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EMBO J. 3:3257-3262(1984).
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                                                                                                                         1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBO J. 4:1251-1255(1985).
                                                                      18; Conservative
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         Query Match
Best Local Similarity
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1 PPLSQETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE=95292092; PubMed=7773777;
Lee W., Harvey T.S., Yin Y., Yau P., Litchfield D., Arrowsmith C.H.,
"Solution structure of the tetrameric minimum transforming domain of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRUCTURE BY NMR OF 319-360.
MEDLINE=94294808; PubMed=8023159;
Clore G.M., Omichinski J.G., Sakaguchi K., Zambrano N., Sakamoto H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDLINE=96197773, PubMed=8632915, Abave P., Freeman J., Bovey R., Iggo R.; Freeman J., Bovey R., Iggo R.; Regulation of specific DNA binding by p53: evidence for a role for 0-g1ycosylation and charged residues at the carboxy-terminus."; Oncogene 12:921-930(1996).
                                                                                                                                                                                                                                                                Anderson C.W., Kieleczawa J., Allalunis-Turner J.;
"Human p53 from the malignant glioma-derived cell lines M059J and M059K have a cancer-associated mutation in exon 8.";
Submitted (MAR-1999) to the EMBL/GenBank/DDBJ databases.
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"High-resolution structure of the oligomerization domain of p53 by
multidimensional NMR.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Addison C., Jenkins J.R., Sturzbecher H.-W.;
"The p53 nuclear localisation signal is structurally linked to a p34cdc2 kinase motif.";
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Bischoff J.R., Friedman P.N., Marshak D.R., Prives C., Beach D
"Human p53 is phosphorylated by p60-cdc2 and cyclin B-cdc2.";
Proc. Natl. Acad. Sci. U.S.A. 87:4766-4770(1990).
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                                                                 Chumakov P.M., Almazov V.P., Jenkins J.R.;
Submitted (JUN-1991) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                  Rozemuller E.H., Tilanus M.G.J.;
"P53 genomic sequence. Corrections and polymorphism.";
Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
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Nat. Struct. Biol. 1:877-890(1994).
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    EMBO J. 10:2879~2887 (1991),
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MINIMAL REPRESSION DOMAIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oncogene 5:423-426(1990).
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                                          SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              normal cells.";
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                         MEDLINE=98026899; PubMed=3321402; MCCOy M., Stavridi E.S., Waterman J.L., Wieczorek A.M., Opella S.J., Halazonetis T.D.; "Hydrophobic side-chain size is a determinant of the three-dimensional structure of the p53 oligomerization domain.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Kussie P.H., Gorina S., Marechal V., Elenbaas B., Moreau J.,
Levine A.J., Pavletich N.P.,
"Structure of the MDM2 oncoprotein bound to the p53 tumor suppressor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=96271983; PubMed=8829653; de Vries T.N., Hartmann A., Blaszyk H., Liao D., Soussi T., Kovach J.S., Sommer S.S.; "Database of mutations in the p53 and APC tumor suppressor genes designed to facilitate molecular epidemiological analyses."; Hum. Mutat. 7:202-213(1996).
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MEDLINE-97081050; PubMed=8875929;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                X-RAY CRYSTALLOGRAPHY (1.7 ANGSTROMS) OF 325-356.
MEDLINE=95184011; PubMed=7878469;
Jeffrey P.D., Gorina S., Pavletich N.P.;
"Crystal structure of the tetramerization domain of the p53 tumor
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                                                                                                                                                                                                                                                                                                     MEDLINE=94294806; PubMed=8023157; Cho Y., Gorina S., Jeffrey P.D., Pavletich N.P.; Cho Y., Gorina S., Jeffrey P.D., Pavletich N.P.; "Crystal structure of a p53 tumor suppressor-DNA complex: understanding tumorigenic mutations."; Science 265:346-355(19494).
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MEDILME=91289166; PubMed=1905840;
Hoolstein M., Sidransky D., Vogelstein B., Harris C.C.;
"p53 mutations in human cancers.";
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Local Similarity 100.0%; Pred. No. 7.3e-08; les 18; Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          suppressor at 1.7 angstroms.";
Science 267:1498-1502(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Genet. 86:369-370(1991).
STRUCTURE BY NMR OF 326-354.
                                                                                                                                                                                                           EMBO J. 16:6230-6236(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Science 262:1980-1981(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Science 274:1001-1005(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Science 274:948-953(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          transactivation domain.'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Harris C.C.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        population.",
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Kham M.A. Hansen C., Welsh J.A., Bennett W.P.;
Kubm M.A., Hansen C., Welsh J.A., Bennett W.P.;
Submitted (FEB-1996) to the EMBL/GenBank/DDBJ databases.
Submitted (FEB-1996) to the EMBL/GenBank/DDBJ databases.
In FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BCI-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUBUNIT: Binds DNA as a homotetramer (By similarity).
SUBCELLULAR LOCATION: Nuclear.
DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                       Macaca fascicularis (Crab eating macaque) (Cynomolgus monkey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EDEEBA1BD3AEAFB8 CRC64;
                                                             15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
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ive 0; Mismatches
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PRINTS; PR00386; P53SUPPRESSR.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      nterPro; IPR002117; P53.
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                                                                                                                                                                                                                                                                                                                                    Cercopithecinae; Macaca.
   STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     325
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                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9541;
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                                                                                                                                                                                                  TP53 OR P53.
P53 MACFA
P56423;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OLIGOMERIZATION.
BASIC (REPRESSION OF DNA-BINDING).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nuclear protein; Phosphorylation; Apoptosis.
DOMAIN
1 44 TRANSCRIPTION ACTIVATION (ACIDIC)
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94.3%; Score 99; DB 1; Length 393; 100.0%; Pred. No. 7.3e-08;
                         0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Khan M.A., Hansen C., Welsh J.A., Bennett W.P.;
Khan M.A., Hansen C., welsh J.A., Bennett W.P.;
Khan M.A., Hansen C. to the EMBL/GenBank/DDBJ databases.

Submitted (FEB-1996) to the EMBL/GenBank/DDBJ databases.

FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of Bcl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-Blood;
MEDLINE-9411042; PubMed-8125305;
KARY H.D., Mountjoy C.P., Wu G., Cornish K.G., Smith L.J.;
"Sequence of a cDNA encoding the p53 protein in rhesus monkey (Macaca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SÜBÜNIT: Binds DNA as a homotetramer (By similarity).
SUBCELLÜLAR LOCATION: Nuclear.
DISBASE: PS3 IS FOUND IN INTREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. PS3 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                             Macaca mulatta (Rhesus macaque).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Cercopithecidae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Transcription regulation; Activator;
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Anti-oncogene; DNA-binding; Transcription regument.

Nuclear protein; Phosphorylation; Apptosis.

44 TRANSCRIPTION ACTIVATION (ACIDIC).
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                                                                                                        Cellular tumor antigen p53 (Tumor suppressor p53).
TP53 OR P53.
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OLIGOMERIZATION.
393 AA
                                                                                          (Rel. 36, Last sequence update) (Rel. 41, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SIMILARITY: BELONGS TO THE P53 FAMILY.
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ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
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                                                               15-JUL-1998 (Rel. 36, Created)
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EMBL; U48956; AAB91534.1; -.
HSSP; P04637; lOLG.
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Best Local Similarity 100.0
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Pfam; PF00870; P53; 1.
   STANDARD;
                                                                                                                                                                                                                                                                                                                           Cercopithecinae, Macaca.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    mulatta).";
Gene 138:223-226(1994).
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                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9544;
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                                                                                                 15-JUL-1998
15-JUN-2002
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Park U., Lee Y.;
"Wild-type p53 sequence of tree shrews.";
"Wild-type p53 sequence of tree shrews.";
Submitted (AuG-1999) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: Acts as tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of Bc1-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            expression.
--- SUBUNIT: Binds DNA as a homotetramer (By similarity).
---- SUBCELLULAR LOCATION: Nuclear.
--- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
IN MANY TYPES OF CANCER.
                                                                                                                                                                                                                                                                                                                                                        Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Scandentia, Tupaiidae, Tupaia.
                                                                                                                                                                                                           15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OLIGOMERIZATION.
                                                                                                                                                                                 Last sequence update)
                                                                                                                                                                                                                                                                                                               Tupaia glis belangeri (Common tree shrew).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -!- SIMILARITY: BELONGS TO THE P53 FAMILY.
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                                 TUPGB SICH. (1997A1) 16-0CT-2001 (Rel. 40, Last seque 16-0CT-2001 (Rel. 41, Last annot 16-0CT (Rel. 41, Last annot 16-0CT (Rel. 41, Last annot 16-0CT (Rel. 41, Last annot 16-0CT)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STRAIN=Chinensis; TISSUE=Liver;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AF175893; AAF22640.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  HSSP; P04637; IYCS.
InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    393 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9396;
                                                                                                                                                                                                                                                                                      OR P53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DNA BIND
DOMAIN
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MOD RES
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RESULT 8
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BASIC (REPRESSION OF DNA-BINDING).
NUCLERRY LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
FD9F36603945A1FA CRC64,
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                              Nuclear protein; Phosphorylation; Apoptosis.

1 44 TRANSCRIPTION ACTIVATION (ACIDIC).
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Gaps
                        .
0
 Length 393;
                        0; Indels
Score 99; DB 1; L. Pred. No. 7.3e-08;
                        Mismatches
94.3.,
100.0%; Fi
         Best Local Similarity 100.
Matches 18; Conservative
Query Match
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0

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0

RESULT 10 P53 BOVIN

1 PPLSQETFSDLWKLLPEN 18 ð

RESULT 9

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                                                                                                                                                                                                                                                                                                                                     PUNA Seq. 5:255-259(1995).

-I- FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BCl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           BASIC (REPRESSION OF DNA-BINDING),
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL),
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                            Dequiedt F., Kettmann R., Burny A., Willems L.;
"Nucleotide sequence of the ovine P53 tumor-suppressor cDNA and its
genomic organization."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -!- SUBUNIT: BINDS DAM AD ...-
-!- SUBCELLULAR LOCATION: Nuclear.
-!- DISEASE: P33 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ProDom: PROJUGE; PS3; 1.
PROSIME; P800248; PS3; 1.
PROSIME; P800348; PS3; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
Nuclear protein; Phosphorylation; Apoptosis.
TRANSCRIPTION ACTIVATION (ACIDIC).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ..
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      expression. SUBUNIT: Binds DNA as a homotetramer (By similarity).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4B024076C79C3B2D CRC64;
                                                   01-OCT-1996 (Rel. 34, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    89.5%; Score 94; DB 1; I 94.4%; Pred. No. 4.2e-07;
                                                                                                                                                                                                                                                                                        Willems L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OLIGOMERIZATION.
 382 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Mismatches
                                                                                                                                                                                                                                                                       MEDLINE=95352828; PubMed=7626788;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
                                   01-OCT-1996 (Rel. 34, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                42809 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; X81705; CAA57349.1; -. HSSP; P04637; 1C26.
                                                                                                                                                                                Caprinae; Ovis.
 STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        17; Conservative
                                                                                                                                                                Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  382 AA;
                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                           Ovis aries (Sheep)
                                                                                                                                                                                                 NCBI_TaxID=9940;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            314
357
300
381
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P53_SHEEP
P51664;
                                                                                                                                                                                                                                                                                        Dequiedt F
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DNA BIND
DOMAIN
                                                                                                                                                                                  Bovidae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.

SPECIES-B.indious; STRAIN-Boran; TISSUE-Blood;
Bishop R.R.P., Gobright B.E.I.;
Bishop R.R.P., Gobright B.E.I.;
Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases.

-I-FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and call type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BCI-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Komori H., Ishiguro N., Horiuchi M., Shinagawa M., Aida Y.,
"Predominant p53 mutations in enzootic bovine leukemic cell lines.",
Vet. Immunol. Immunopathol. 52:53-63(1996).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUBUNIT: Binds DNA as a homotetramer (By similarity).
SUBCELLULAR LOCATION: Nuclear.
DISEASE: P53 IS FOUND IN INVERASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
IN MANY TYPES OF CANCER.
                                                                                                                                                         Bos indicus (Zebu).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anti-oncogene; DNA-binding; Transcription regulation; Activator; Nuclear protein; Phosphorylation; Apoptosis.
                                                                                                                                                                                                                                                                                            SPECIES-Bovine; TISSUE-Liver;
MEDLINES-9535.28423; PubMed=7626789;
Dequiedt F., Kettmann R., Burny A., Willems L.;
"Nucleotide sequence of the bovine P53 tumor-suppressor CDNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TRANSCRIPTION ACTIVATION (ACIDIC)
                                                                                                Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE OF 13-386 FROM N.A.
SPECIES=Bovine; STRAIN=Holstein; TISSUE=Thymus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OLIGOMERIZATION.
                                    OJSOCA, 1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BY SIMILARIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=96401400; PubMed=8807776;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL, X81704, CAA57348.1; -. EMBL, D49825; BAA08629.1; -. EMBL, U74486; AAB51214.1; -. HSSP; P04637; ITUP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                       DNA Seq. 5:261-264(1995).
                                                                                                                                         Bos taurus (Bovine), and
        STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             285
                                                                                                                                                                                                             Bovidae, Bovinae, Bos.
NCBI_TaxID=9913, 9915;
                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DNA BIND
DOMĀIN
P53 BOVIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DOMAIN
MOD_RES
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NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinasses. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of BGl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDLINE-97376996; PubMed=9233767; Feitelson M.A., Ranganathan P.N., Clayton M.M., Zhang S.M.; Feitelson M.A., Ranganathan P.N., Clayton M.M., Zhang S.M.; "Partial characterization of the woodchuck tumor suppressor, p53, and its interaction with woodchuck hepatitis virus X antigen in Oncogene 15:327-336(1997).
                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DISEASE: P3 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                                                                                                                                                                                                                Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Rodentia, Sciurognathi, Sciuridae, Sciurinae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Nuclear protein; Phosphorylation; Apoptosis.

DOMAIN 1 44 TRANSCRIPTION ACTIVATION (ACIDIC).

DNA_BIND 100 259 BY SIMILARITY.

DOMAIN 323 354 OLIGOMERIZATION.
                                                                                           ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBUNIT: Binds DNA as a homotetramer (By similarity). SUBCELLULAR LOCATION: Nuclear.
                                                       Length 386;
                                                                                           1; Indels
380 380 R -> T (IN REF. 2).
386 AA; 43255 MW; 222473F28C548F31 CRC64;
                                                                                                                                                                                                                                                                                                                                         15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                           Pred. No. 4.3e-07;
                                                       89.5%; Score 94; DB 1; 94.4%; Pred. No. 4.3e-07
                                                                                                                                                                                                                                                                 PRT; 391 AA
                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                           Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; AJ001022; CAA04478.1; -. HSSP; P04637; 1TUP.
InterPro; IPR002117; P53.
Pfam; PR00870; P53; 1.
PRINTS; PR00386; P535UPPRESSR.
ProDom; P0002681; P53; 1.
                                                                                                                                                                                                                                                                                                         16-OCT-2001 (Rel. 40, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                IN MANY TYPES OF CANCER.
                                                                                                                                                                                                                                                                                                                         (Rel. 40, Last (Rel. 41, Last
                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                          12 PPLSQETFSDLWNLLPEN 29
                                                                                               17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                    Marmota monax (Woodchuck)
                                                                                                                                                                                                                                                                     STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    354
385
321
390
                                                           Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         expression.
                                                                                                                                                                                                                                                                                                                               L6-OCT-2001
                                                                                                                                                                                                                                                                     P53 MARMO
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MOD RES
                       SEQUENCE
     CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Na rmot a
                                                                                                                                                                                                                                                                                       036006;
                                                                                                   Matches
                                                                                                                                                                                                                                  RESULT 11
P53_MARMO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Setoguchi A., Sakai T., Okuda M., Minehata K., Yazawa M., Ishizaka T.,
Watari T., Hasagawa A., Tsujimoto H.,
"Aberrations of p53 tumor suppressor gene in various spontaneous
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=9532915; pubMed=7600529; Kraegel S.A., Pazzi K.A., Madewell B.R.; Fraegel S.A., Pazzi K.A., Madewell B.R.; Sequence analysis of Ganine p53 in the region of exons 3-8."; Cancer Lett. 92:181-186(1995).

--- FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinasses. Apoptosis induction seems to be mediated either by stimulation of BAZ and FAS antigen expression, or by repression of Bc1-2
                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE=Leukocyte;
MEDLINE=288178696; PubMed=9519881;
Veldhoen N., Milner J.;
"Isolation of canine p53 cDNA and detailed characterization of the full length canine p53 protein.";
Oncogene 16:1077-1084(1998).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    -!- SUBUNIT: Binds DNA as a homotetramer (By similarity).
-!- SUBCELLULAR LOCATION: Nuclear.
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Canis familiaris (Dog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Carnivora, Fissipedia, Canidae, Canis.
                                                                                                                                             ..
                                                        89.5%; Score 94; DB 1; Length 391; 94.4%; Pred. No. 4.3e-07; Live 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        tumors in the dog.";
Submitted (DEC-1998) to the EMBL/GenBank/DDBJ databases.
391 AA; 43468 MW; EIDESDB84BA40182 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               15-JUN-2002 (Rel. 41, Last annotation update)
Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                            P53_CANFA STANDARD, PRT; 381 AA. Q29537; Q9TV78; 01-NOV-1997 (Rel. 35, Created) 15-DEC-1998 (Rel. 37, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE OF 25-300 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 IN MANY TYPES OF CANCER.
                                                                                                                                                                                              1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                     12 PPLSQETFSDLWNLLPEN 29
                                                                                                                                      Conservative
                                                        Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9615;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FISSUE=Spleen;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRAIN-Beagle;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TP53 OR P53
                                                                                                                                      17;
SEOUENCE
                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                             RESULT 12
                                                                                                                                                                                                                                                                                                                                                                                                P53_CANFA
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EMBL; AF060514; AAC16909.1; -. EMBL; AR020761; BAA78379.1; -.

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BY SIMILARITY.
OLIGOMERIZATION.
BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYATION (BY SIMILARITY).
MEES -> MOEP (IN REF. 2).
L -> P (IN REF. 2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  J. Vet. Med. Sci. 55:801-805(1993).

-! FUNTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of Bc1-2
                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=94114699; PubMed=8286534; Okuda M., Umeda A., Matsumoto Y., Momoi Y., Watari T., Goitsuka R., O'Erien S.J., Tsujimoto H., Hasegawa A.; "Molecular cloning and chromosomal mapping of feline p53 tumor suppressor gene.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUB=Lymph node;

MEDLINE=5433960; PubMed=8056458;

Okuda M., Umeda A., Sakai T., Ohashi T., Momoi Y., Youn H.Y.,

Matari T., Goltsuka R., Tsujimoto H., Hasegawa A.;

"Cloning of feline p53 tumor-suppressor gene and its aberration in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   -!- SUBUNIT: Binds DNA as a homotetramer (By similarity).
-!- SUBCELULAR LOCATION: Nuclear.
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Felis silvestris catus (Cat).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Carnivora, Fissipedia, Felidae, Felis.
                                                                                                              Anti-oncogene; DNA-binding; Transcription regulation; Activator; Nuclear protein; Phosphorylation; Apoptosis.

1 44 TRANSCRIPTION ACTIVATION (ACIDIC).
                                                                                                                                                                                                                                                                                                                                                        ·.
0
                                                                                                                                                                                                                                                                                                                       Length 381;
                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                761A718FDC93DA59 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                   Score 90; DB 1; I
Pred. No. 1.7e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      386 AA.
                                                                                                                                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   IN MANY TYPES OF CANCER.
SIMILARITY: BELONGS TO THE P53 FAMILY.
               HSSP, P04637; 10LG.
InterPro; IPR002117; P53.
Pffan, PF00870; P53; 1.
PRINTS; PR00386; P533UPPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 int. J. Cancer 58:602-607(1994).
                                                                                                                                                                                                                                                                                     42486 MW;
EMBL; S77819; AAB42022.1; -.
                                                                                                                                                                                                                                                                                                                   85.78;
                                                                                                                                                                                                                                                                                                                                   88.98;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE OF 34-354 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                  1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                12 PPLSQETFSELWNLLPEN 29
                                                                                                                                                                                                                                                                                                                                                     16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STANDARD;
                                                                                                                                                                                 344
                                                                                                                                                                                                                                 380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hematopoietic tumors
                                                                                                                                                                                                                                                                                  381 AA;
                                                                                                                                                                                                                                                                                                                                   Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
                                                                                                                                                                                 313
356
299
380
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID=9685;
                                                                                                                                                                   83
                                                                                                                                                                 DNA BIND
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P53 FELCA
                                                                                                                                                                                                                                           CONFLICT
                                                                                                                                                                                                                                                                                  SEQUENCE
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                                                                                                                                                                                                 DOMAIN
                                                                                                                                                                                                                  DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P41685;
                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 13
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TISSUE=Embryonic fibroblast;
Shimizu T., Nikaido O., Suzuki F.;
Submitted (JUN-1996) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases.
                                                                                                                                                                                                                                                                                                                                                                       BASIC (REPRESSION OF DNA-BINDING).
NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Rodentia, Sciurognathi, Muridae, Cricetinae,
                                                                                                                                                                                               InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00346; P53SUPPRESSR.
ProDom; P0002661; P53; 1.
PROSITE; P800346; P53; 1.
Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ..
                                                                                                                                                                                                                                                                                                      Nuclear protein, Phosphorylation, Apoptosis.

1 44 TRANSCRIPTION ACTIVATION (ACIDIC)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Lee H., Larner J.M., Hamlin J.L.; "Cloning and characterization of Chinese hamster p53 cDNA.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              85.7%; Score 90; DB 1; Length 386;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chaung W., Mi L.J., Boorstein R.J.;
Submitted (MAR-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                           42692 MW; D08B43BA1BC8EB78 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             009183, Q64397, P97258, P97788, 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update) 15-JUN-2002 (Rel. 41, Last annotation update) 17-F3 OR P53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Pred. No. 1.8e-06;
                                                                                                                                                                                                                                                                                                                                       BY SIMILARITY.
OLIGOMERIZATION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     393 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                            K -> R (IN REF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cricetulus griseus (Chinese hamster).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=97183659; PubMed=9031625;
                                                                                                                                               EMBL; D26608; BAA05653.1; -. EMBL; D16460; BAA03927.1; -. HSSP; P04637; IOLG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   12 PPLSQETFSELWNLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gene 184:177-183(1997).
                                                                                                                                                                                                                                                                                                                                                       349
                                                                                                                                                                                                                                                                                                                                                                                                                                           386 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=10029;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cricetulus.
                                                                                                                                                                                                                                                                                                                      DOMAIN
DNA BIND
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     P53 CRIGR
                                                                                                                                                                                                                                                                                                                                                                                                         MOD RES
CONFLICT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                           DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 14
P53_CRIGR
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MEDLINE=92210007; PubMed=155573;
Legros Y., McIntyre P., Soussi T.;
"The cDNA cloning and immunological characterization of hamster p53.";
Gene 112:247-250(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BY SIMILARITY.

OLIGOMERIZATION.

DASJIC (REPRESSION OF DNA-BINDING).

NUCLEAR LOCALIZATION SIGNAL (POTENTIAL).

PHOSPHORYLATION (BY SIMILARITY).

L -> Q (IN CELL LINE V79-4).

C -> W (IN CELL LINE V79-4).

Y -> F (IN REF. 2).

WW; 2A7830E788311689 CRC64;
Apoptosis induction seems to be mediated either by stimulation of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                      SUBCELLULAR LOCATION: Nuclear.
SUBCELLULAR LOCATION: Nuclear Location and Language Location and Location and Language Loca
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Butheria, Rodentia, Sciurognathi, Muridae, Cricetinae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anti-oncogene; DNA-binding; Transcription regulation; Activator;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nuclear protein; Phosphorylation; Apoptosis.

OMAIN

1 45
TRANSCRIPTION ACTIVATION (ACIDIC)
                           BAX and FAS antigen expression, or by repression of Bcl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     81.9%; Score 86; DB 1; Length 393; 94.1%; Pred. No. 7.4e-06;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   i; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cellular tumor antigen p53 (Tumor suppressor p53).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P53 MESAU STANDARD, PRT; 396 AA. 000366; P97276; 01-DEC-1992 (Rel. 24, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                               SIMILARITY: BELONGS TO THE P53 FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mesocricetus auratus (Golden hamster).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00386; P538UPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      43378 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; Y08900; CAA70108.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; U50395; AAC53040.1; -. EMBL; D86070; BAA13004.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2 PLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13 PLSQETFSDLWKLLPPN 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       133
135
103
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356
387
323
392
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325
368
311
392
133
103
393 AA;
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SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DNA BIND
DOMAIN
DOMAIN
DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 15
P53_MESAU
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Hou E.W., Wiseman R.;
Submitted (APR-1994) to the EMBL/GenBank/DDBJ databases.
-! FUNCTION: Acts as a tumor suppressor in many tumor types; induces growth arrest or apoptosis depending on the physiological circumstances and cell type. Involved in cell cycle regulation as a trans-activator that acts to negatively regulate cell division by controlling a set of genes required for this process. One of the activated genes is an inhibitor of cyclin-dependent kinases. Apoptosis induction seems to be mediated either by stimulation of BAX and FAS antigen expression, or by repression of Bcl-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         BY SIMILARITY.
OLIGOMERIZATION.
DASJIC (FEPRESSION OF DNA-BINDING).
NUCLERA LOCALIZATION SIGNAL (POTENTIAL).
PHOSPHORYLATION (BY SIMILARITY).
                                                                                                                                                                                 expression.
-!- SUBUNIT: Binds DNA as a homotetramer (By similarity).
-!- SUBCELLULAR LOCATION: Nuclear.
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY
OF TRANSPORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED
IN MANY TYPES OF CANCER.
-!- SIMILARITY: BELONGS TO THE P53 PAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; M75144; AAA37085.1; -. BMBL; U07182; AAB41344.1; -. PIR; JH0633; JH0633. HSSP; P04637; 1TUP: InterPer; IRFO0870; P53; 1. PFAM; PFO0870; P53; 1. PRINTS; PRO0386; P53SUPPRESSR.
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395
395
188
396 AA;
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DNA BIND
DOMAIN
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Search completed: February 12, 2003, 10:35:02 Job time : 12 secs

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0; Gaps

81.9%; Score 86; DB 1; Length 396; 94.1%; Pred. No. 7.5e-06; ive 0; Mismatches 1; Indels

ilarity 94.1%; Conservative

Best Local Similarity Matches 16; Conserv

Query Match

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G -> S (IN REF. 2). 906EF02568099BE3 CRC64;

43631 MW;

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February 12, 2003, 10:33:27; Search time 29 Seconds (without alignments) 134.996 Million cell updates/sec
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GenCore version 5.1.3 Copyright (c) 1993 - 2003 Compugen Ltd.
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                                                                                                                                                                                                                                                                                                                   671580 segs, 206047115 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                      OM protein - protein search, using sw model
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105
1 PPLSQETFSDLWKLLPENG 19
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sp bacteria: *
sp bungi: *
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

sp_unclassified:*

111: 12: 13: 15: 17:

sp_rvirus:*
sp_bacteriap:*

sp_archeap:*

sp_virus:*
sp_vertebrate:*

sp_rodent:* sp_virus:* sp plant: *

SS		Description	Q9nzd0 homo sapien	Q8spz3 delphinapte	Oguq61 homo sapien	homo	homo	homo	homo	homo		homo	homo	Q16810 homo sapien	Q9btm4 homo sapien	Q16535 homo sapien	O9n252 sus scrofa	Q920y0 meriones un
SUMMARIES		ID	002N60	Q8SPZ3	090061	Q9HAQ8	016811	016807	015086	Q15087	Q1508B	016808	Q16809	016810	Q9BTM4	Q16535	Q9N252	Q920Y0
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		Match Length DB	32	387	393	393	393	393	393	393	393	393	393	393	393	393	387	390
	* Query	Match	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	94.3	89.5	83.8
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12 PPLSQETFSDLWKLLPEN 29

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77 73.3 25 4 QNRP68 74 70.5 397 11 Q9ER40 75 70.5 397 11 Q9ER40 76 8.6 370 11 Q9ERP9 76 8.6 271 11 Q9ERP9 60 57.1 265 13 Q9W681 60 57.1 265 13 Q9W682 52 49.5 41 265 13 Q9W682 52 49.5 443 5 Q9NGC7 51.5 49.0 1009 11 Q62640 51.5 49.0 1009 11 Q61427 60 47.6 47.1 Q9TH30 60 47.6 47.1 Q9TH30 61 47 44.8 157 12 Q91H93 61 44.8 157 12 Q91H93 61 44.8 158 12 Q91H32 61 69H0K7	Д.	O9eq10 rattus norv P89002 pracomys nat Q9w680 oncorhynchu Q9w681 oncorhynchu Q9w682 oncorhynchu Q9qqq2 rhizobium m Q9qqqq mya arenari	Q911k0 homo sapien Q62640 rattus norv Q61627 mus musculu Q9chk4 lactococcus Q9xiq5 arabidopsis Q8ra97 thermoanaer Q8try 0 methanopyru Q8t4n3 rhipicephal Q8yjk2 brucella me	Q817v3 spisula sol Q91u98 pepper mild Q91u97 pepper mild Q91e32 pepper mild Q83483 tobacco mos Q9sjm2 arabidopsis Q9huk7 pseudomonas
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ALIGNMENTS

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TISSUESTITE FROM N.A.

TISSUE-GLIOMAS AND ASSOCIATED NORMAL TISSUES;

TISSUE-GLIOMAS AND ASSOCIATED NORMAL TISSUES;

TISSUE-GLIOMAS AND ASSOCIATED NORMAL TISSUES;

Thompson-Hehir J., Davises M.P.A., Green J.A., Halliwell N.,

JOYCE K.A., Salisbury J., Sibson D.R., Vergote I., Walker C.;

"Mutation Detection Utilizing a Novel PCR Approach for Amplification of the p53 dene from Microdissected Tissue: Application to Archival TL Tumor Samples.";

Submitted (DEC-1999) to the EMBL/GenBank/DDBJ databases.

SERIJ, AF2103009; AAF63442.1; -.

REMBL, AF2103009; AAF63442.1; -.

REMBL, AF2103009; AAF63442.1; JOINED.

INTERPORT JPRO02117; P53.

Probom; PD002611; P53; 1.

Probom; PD002611; P53; 1.

PRODOM; PN0VER 32 AA; 3670 MW; FOROEA983A90B917 CRC64;
                                                                                                                                                                          Homo sapiens (Human).
Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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                                                                            01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
                                         32 AA.
                                                                                                                                       P53 tumor suppressor protein (Fragment). TP53.
                                         PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100.
Matches 18; Conservative
                                         PRELIMINARY;
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RESULT 1
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Pfeiffer C.J., Sato S.;
"Nucleotide sequence of the p53 cDNA of beluga whale (Delphinapterus
                                                                                                                                                                                                                                         0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Anderson C.W., Kieleczawa J., Allalunis-Turner M.J.;
"Human p53 from malignant glioma-derived M059J and M059K cells has a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Isolation of two cell lines from a human malignant glioma specimen differing in sensitivity to radiation and chemotherapeutic drugs."; Radiat. Res. 134:349-354(1993).
                                                                                     Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Butheria, Cetartiodactyla, Cetacea, Odontoceti,
Monodontidae, Delphinapterus.
                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Allalunis-Turner M.J., Barron G.M., Day R.S. III, Dobler K.D., Mirzayans R.;
                                                                                                                                                                                                                      Length 387;
                                                                                                                                                                                                             Score 99; DB 6; Length 30;
Pred. No. 9e-08;
                                                                                                                                                                                          EMBL; AF475081; AAL83290.1; -.
SEQUENCE 387 AA; 43033 MW; E4C8BEDBF34A540E CRC64;
                                 01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                        01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                                                                                                                                         PRT; 393 AA.
                                                                                                                                                                                                                    Query Match

94.3%; Score 99; DB
Best Local Similarity 100.0%; Pred. No. 9e-
Matches 18; Conservative 0; Mismatches
                                                                               Delphinapterus leucas (Beluga whale)
                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDLINE=93303270; PubMed=8316628;
                                                                                                                                                                                                                                                         1 PPLSOETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                    12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                                                                                                                                                                                     Cellular tumor antigen p53.
                PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (Human)
                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
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                                                                                                                                                                                Gene 0:0-0(2002).
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                                                                                                                                                                                                   SEQUENCE
                           Q8SPZ3;
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                 O8SPZ3
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RESULT 2
          Q8SPZ3
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"Hyaluronidase Induction of a WW Domain-Containing Oxidoreductase that
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J. Blol., Chem. 276;13361-3370(2001).

-!- FUNCTION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES GROWTH ARREST OR APOPTOSIS DEPENDING ON THE PHYSIOLOGICAL CIRCUMSTANCES OR CELL TYPE, BUT BOTH ACTIVITIES ARE INVOLVED IN TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATION, IS A TRANS-ACTIVATOR THAT ACTS TO NEGATIVELY REGULATE CELL DIVISION BY CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES. APOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRINTS; PR00386; P53SUPPRESSR.
ProDom; P0020881; P53; 1.
PROSITE; PS00348; P53; 1.
Activator; Anti-oncogene; Apoptosis; DNA-binding; Nuclear protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
MEDLINE=21264809; PubMed=11058590;
Chang N.-S., Pratt N., Heath J., Schultz L., Sleve D., Carey G.B.,
Zevotek N.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                   Score 99; DB 4; Length 393;
Pred. No. 9.2e-08;
                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
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SEQUENCE 393 AA; 43712 MW; AC611E4938C7BC3B CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         94.3%; Scor.
100.0%; Pred. No. >--.
MANY TYPES OF CANCER (BY SIMILARITY).

-!- SIMILARITY: BELONGS TO THE P53 FAMILY.
EMBL; AF135121; AAD28535.1; --
EMBL; AR135120; AAD28535.1; JOINED.
HSSP; P04637; 10LG.
INTERPO: IPTRO02117; P53.
FAMILY PRO03107; P53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT; 393 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                 ProDom; PD002681; P53; 1. PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 100.0
Matches 18; Conservative
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12 PPLSQETFSDLWKLLPEN 29

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MEDLINE=87064416; PubMed=2946935;

Lamb P., Crawford L.;

Lamb P., Crawford L.;

Caracterization of the human p53 gene.";

Mol. Cell. Biol. 6:1379-1385(1986).

-!- FUNCTION: ACTS AS A TUMOR SUPPRESSON IN MANY TUMOR TYPES. INDUCES GROWTH ARREST OR APOPTOSIS DEPENDING ON THE PHYSTOLOGICAL CIRCUMSTANCES OR CELL TYPE, BUT BOTH ACTIVITIES ARE INVOLVED IN TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATION, IS A TRANS-ACTIVATED GENES IS TO NEGATIVELY REGULATION, IS A TRANS-CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.

APOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF BAX AND FAS ANTIGEN EXPRESSION, OR BY REPRESSION OF BCL-2

EXPRESSION (BY SIMILARITY).

C. -1- SUBCELLULAR LOCATION: NUCLEAR (BY SIMILARITY).

C. -1- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF TRANSFORMED CELLS. P53 IS FREQUENTLY MITATED OR INACTIVATED IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Isolation and characterization of a human p53 cDNA clone: expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Activator; Anti-oncogene; Apoptosis; DNA-binding; Nuclear protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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94.3%; Score 99; DB 4; Length 393;
Best Local Similarity 100.0%; Pred. No. 9.2e-08;
Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
MEDLINE-85126934; PubMed-6396087;
Matlashewski G., Lamb P., Pim D., Peacock J., Crawford L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Phosphorylation, Transcription regulation.
SEQUENCE 393 AA, 43698 MW, AD1979573AFA1DA5 CRC64;
                                                                                                                                                                                                                                                                01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
                                                                                                                                                                                                       393 AA.
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M13118; AAA59987.1; JOINED.
M13119; AAA59987.1; JOINED.
M13120; AAA59987.1; JOINED.
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M13112, AAA59987.1; JOINED.
M13113, AAA59987.1; JOINED.
M13114; AAA59987.1; JOINED.
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ProDom; PD002681; P53; 1.
                                                                                                                                                                                                                                                                                                                                                           Cellular tumor antigen p53.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             of the human p53 gene.";
EMBO J. 3:3257-3262(1984).
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Pfam; PF00870; P53; 1.
1 PPLSORTFSDLWKLLPEN 18
                                                     12 PPLSQETFSDLWKLLPEN 29
                                                                                                                                                                                                          PRELIMINARY;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
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                                                                                                                                                                                                                                         016811;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL;
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EMBL;
EMBL;
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EMBL;
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                                                                                                                                                 RESULT 5
                                                                                                                                                                                 01681
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LEMBO J. 10.2879-2887 (1991).

CIRCHORTION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES C. I- FUNCTION: ACTS AS A TUMOR SUPPRESSOR IN THE PHYSIOLOGICAL GROWTH ARREST OR APPOPTOSIS DEPENDING ON THE PHYSIOLOGICAL INVOLVED IN CIRCHARTANCES OR CELL TYPE, BUT BOTH ACTIVITIES ARE INVOLVED IN TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATE CELL DIVISION BY ACTIVATOR THAT ACTS TO NEGATIVELY REGULATE CELL DIVISION BY CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASS. APPOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF BAX AND FAS ANTIGEN EXPRESSION, OR BY REPRESSION OF BCL-2

EXPRESSION (BY SIMILARITY).

-!- SUBCELLULAR LOCATION: NUCLEAR (BY SIMILARITY).

-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF TRANSFORMED CELLS. PS3 IS FREQUENTLY MITATED OR INACTIVATED IN MANY TYPES OF CANCER (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; X60011; CAA42626.1; -.
HSSP; P04637; 10LG.
HTAETPC: IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
PRODOM; P0002861; P53; 1.
PROSITE; PS00148; P53; 1.
Activator; Anti-oncogene; Apoptosis; DNA-binding; Nuclear protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                               Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
MEDIATE=29.007731; PubMed=1915267;
Farrell P.J., Allan G., Shanahan F., Vousden K.H., Crook T.;
Farrell F.J., Allan G., Shanahan F., Vousden K.H., Erook T.;
Farrell F.J., Allan G., Shanahan F., Vousden K.H., Erook T.;
Farrell F.J., Allan G., Shanahan F., Vousden K.H., Erook T.;
Farrell F.J., Allan G., Shanahan F., Vousden K.H., Crook T.;
Farrell F.J., Allan G., Shanahan F., Vousden K.H., Crook T.;
Farrell F.J., Allan G., Shanahan F., Vousden K.H., Crook T.;
Farrell F.J., Farrell F.J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94.3%; Score 99; DB 4; Length 393;
100.0%; Pred. No. 9.2e-08;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      393 AA; 43731 MW; 3A3580718E932E27 CRC64;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                     Last sequence update)
Last annotation update)
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VARIANT 193 193 R -> H.
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                                                      393 AA
                                                                                                                                                                                                                             Cellular tumor antigen p53 (Fragment)
                                                                                                                          Created)
                                                      PRT;
                                                                                                                       01-NOV-1996 (TrEMBLrel. 01, 01-NOV-1996 (TrEMBLrel. 01, 01-MAR-2002 (TrEMBLrel. 20,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 12 PPLSQETFSDLWKLLPEN 29
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                                                             PRELIMINARY;
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9606;
                                                                                                                                                                                                                                                            TP53 OR P53
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SEQUENCE
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RESULT 6
                                  016807
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RN GEOURNEE FROM N.A.

RN SEQUENCE FROM N.A.

RA FAITELINE-9200731; PubMed=1915267;

RA FAITELINE-9200731; PubMed=1915267;

RA FAITELINE-9200731; PubMed=1915267;

REDIANS-9200731; Pubmmed=1915267;

REDIANS-9200731;

REDIANS-920073
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                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                          SEQUENCE FROM N.A.
MEDLINE=92007731; PubMed=1915267;
Farrell P.J. Allan G., Shanahan F., Vousden K.H., Crook T.;
"p53 is frequently mutated Burkitt's lymphoma cell lines.";
EMBO O. 10:2879-2887(1991).
EMBO D. 16:3879-2887(1991).
EMBL; X60016; CAA42631.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  94.3%; Score 99; DB 4; Length 393; 100.0%; Pred. No. 9.2e-08; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                              393 AA; 43713 MW; AD48ABDB26FA208B CRC64;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
cellular tumor antigen p53 (Fragment).
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                                                                                                                                                                                                                                                                                                        InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
VARIANT 238 238 Y
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Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 PPLSQETFSDLWKLLPEN 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18; Conservative
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SEQUENCE
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
N [1]

P SEQUENCE FROM N.A.

X MEDLINE-2007731; PubMed=1915267;

XA Farrell P.J., Allan G., Shanahan F., Vousden K.H., Crook T.;

RI "p53 is frequently mutated Burkitt's lymphoma cell lines.";

RL BMBO J. 10:2879-2887(1991).

BMBO J. 10:2879-2887(1991).

BR SP; PO4637; IOLG.

BR HSSP; PO4637; IOLG.

BR HSSP; PO4637; IOLG.

BR HSSP; PO003117; P53.

BR TherPro; IPR003117; P53.

BR PRINTS; PR00386; P5331.

BR PRINTS; PR00368; P5331.

PRODOM; P0002681; P53; 1.

WADTANT 246 246 T -> M.
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MEDLIARE=92007731; PubMed=1915267;
MEDLIARE=92007731; PubMed=1915267;
Farrell P.J. Allan G., Shanahan F., Vousden K.H., Crook T.;
"p53 is frequently mutated Burkitt's lymphoma cell lines.";
EMBL Of 10:2879-2887(1991).
EMBL; X60014; CAA42629.1; -.
HSSP; P04637; 10LG.
                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 94.3%; Score 99; DB 4; Length 393; Best Local Similarity 100.0%; Pred. No. 9.2e 08; Matches 18; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      94.3%; Score 99; DB 4; Length 393; 100.0%; Pred. No. 9.2e-08; Ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        393 AA; 43694 MW; C6A118E550EB96BB CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
P53 transformation suppressor (Fragment).
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last_annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT;
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Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
ProDom; PD002681; P53; 1.
VARLANT 237 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
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O15087;
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RESULT 8

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01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
Cellular tumor antigen p53 (Fragment).
TP53 OR P53.
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                                                                Homo sapiens (Human).
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TISSUE=KIDNEY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9606;
                                                                                                                  NCBI_TaxID=9606;
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NON TER
SEQUENCE
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CHONCITION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES GROWTH ARREST OR APOPTOSIES DEPENDING ON THE PHYSICLOGICAL CIRCUMSTANCES OR CELL TYPE, BUT BOTH ACTIVITIES ARE INVOLVED IN TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATION, IS A TRANSCRIPTOR THAT ACTS TO MEGATIVELY REGULATE CELL DIVISION BY ACTIVATION THAT ACTS TO MEGATIVELY REGULATE CELL DIVISION BY CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES. APOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF BAX AND FAS ANTIGEN EXPRESSION OR BY REPRESSION OF BCL-2 EXPRESSION OR BY SIMILARITY).

CHOSTOLILAR HOCATION: NUCLEAR (BY SIMILARITY).

CHOSTOLILAR HOCATION: NUCLEAR BY SIMILARITY).

CHOSTOLILAR HOCATION: NEWS SIMILARITY).

CHOSTOLILAR HOCATION: NUCLEAR BY SIMILARITY).

CHOSTOLILAR HOCATION: PS3: 1.

CHOSTOLILAR HOCATION: PS3: 1.
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Phosphorylation; Transcription regulation.
                                                                                                                                                                                                                                                                                                                                                                 Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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MEDLINE-92007731; PubMed=1915267;
MEDLINE-92007731; PubMed=1915267;
Sarrell P., Allan G., Shanahan F., Vousden K.H., Crook T.;
"p53 is frequently mutated Burkitt's lymphoma cell lines.";
EMBO J. 10:2879-2887(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Match 94.3%; Score 99; DB 4; Length 393; Local Similarity 100.0%; Pred. No. 9.2e-08; es 18; Conservative 0; Mismatches 0; Indels
                                                    94.3%; Score 99; DB 4; Length 393; 100.0%; Pred. No. 9.2e-08; tive 0; Mismatches 0; Indels
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                         393 AA; 43627 MW; 3325FAE63F8E1E61 CRC64;
                                                                                                                                                                                                                                                                      Created)
Last sequence update)
Last annotation update)
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Last sequence update)
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                                                                                                                                                                                                                                         393 AA.
                                                                                                                                                                                                                                                                                                                      Cellular tumor antigen p53 (Fragment).
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                                                                                                                                                                                                                                      PRT;
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                                                                                                                                                                                                                                                       Q16809;
01-NOV-1996 (TrEMBLrel. 01,
01-NOV-1996 (TrEMBLrel. 01,
01-MAR-2002 (TrEMBLrel. 20,
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                                                                                                                                             12 PPLSQETFSDLWKLLPEN 29
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                                                                                          18; Conservative
                                                                                                                                                                                                                                         PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                       Homo sapiens (Human)
                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9606;
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01-NOV-1996
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                             SQ SEQUENCE
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                                                               Query Match
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REDIENCE FROM N.A.

MEDLINE=9200731; PubMed=1915267;

MEDLINE=92007731; PubMed=1915267;

MEDLINE=10:2819-2887(1991)

MEDLINE=10:2819-2819

MEDLINE=10:2819

MEDLINE=10
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Submitted (FEB-2001) to the EMBL/GenBank/DDBJ databases.
-!- FUNCTION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES GROWTH ARREST OR APOPTOSIS DEPENDING ON THE PHYSIOLOGICAL CIRCUMSTANCES OR CELL TYPE, BOT BOTH ACTIVITIES ARE INVOLVED IN TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATION, IS A TRANS-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JUN-2001 (TrEMBLrel. 17, Last sequence update) 01-MAR-2002 (TrEMBLrel. 20, Last annotation update)
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PRINTS, PR00386; P5338UPPRESSR.
ProDom; P0002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                94.3%;
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les 18; Conservative
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           CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.
APOPTOSIS INDUCTION SEEMS TO BE MEDIATED EITHER BY STIMULATION OF BAX AND FAS ANTIGEN EXPRESSION, OR BY REPRESSION OF BCL-2
EXPRESSION (BY SIMILARITY)
                                                                                                                                                                                                                                                                                                                                   0; Gaps
                                                                            -!- SUBCELLULAR LOCATION: NUCLEAR (BY SIMILARITY).
-!- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF
TRANSFORMED CELLS. P53 IS FREQUENTLY MUTATED OR INACTIVATED IN
MANY TYPES OF CANCER (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE P53 FAMILY.

EMBL; BC003596; AAH03596.1;
                                                                                                                                                                                                                                          Activator; Anti-oncogene; Apoptosis; DNA-binding; Nuclear protein; Phosphorylation; Transcription regulation.
SEQUENCE 393 AA; 43686 MW; 5C701E458E016756 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
 ACTIVATOR THAT ACTS TO NEGATIVELY REGULATE CELL DIVISION BY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .;
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                                                                                                                                                                                                                                                                                                    94.3%; Score 99; DB 4; Length 393; 100.0%; Pred. No. 9.2e-08; Live 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                  0; Indels
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT; 393 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     transformation suppressor (Fragment).
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                                                                                                                                                                         InterPro, IPR002117; P53.
Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
PRODOM; P0002681; P53; 1.
PROSITE; PS00348; P53; 1.
                                                                                                                                                                                                                                                                                                                                                      1 PPLSOETFSDLWKLLPEN 18
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                                                                                                                                                                                                                                                                                                                              18; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Homo sapiens (Human)
                                                                                                                                                               HSSP; P04637; 10LG.
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                                                                                                                                                                                                                                                                                                            Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9606;
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Q16535
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Plass M., Hartmann E., Mueller O., Kuhlmann J.;

Alfaentification of a Sus scrofa gene encoding a protein with high
similarity to p53 from Homo sapiens, Bos taurus and Ovis aries.";
Submitted (JAN-1999) to the BMEL/GenBank/DDBJ databases.

-!- FUNCTION: ACTS AS A TUMOR SUPPRESSOR IN MANY TUMOR TYPES. INDUCES
GROWTH ARREST OR APOPTOSIS DEPENDING ON THE PHYSTOLOGICAL
CIRCUMSTANCES OR CELL TYPE, BUT BOTH ACTIVITIES ARE INVOLVED IN
TUMOR SUPPRESSION. ACTS IN CELL CYCLE REGULATE CELL DIVISION BY
CONTROLLING A SET OF GENES REQUIRED FOR THIS PROCESS. ONE OF THE
ACTIVATED GENES IS AN INHIBITOR OF CYCLIN-DEPENDENT KINASES.
ACTIVATED GENES IS AN INHIBITOR OF STIMULATION OF BAX AND FAS ANTIGEN EXPRESSION, OR BY REPRESSION OF BCL-2
EXPRESSION (BY SIMILARITY).
--- SUBGELLULAR LOCATION; NUCLEAR (BY SIMILARITY).
--- SUBGELLULAR LOCATION; NUCLEAR (BY SIMILARITY).
--- DISEASE: P53 IS FOUND IN INCREASED AMOUNTS IN A WIDE VARIETY OF MANY TYPES OF CANCER.

MANY TYPES OF CANCER. (BY SIMILARITY).
--- STANT BYTHES OF CANCER (BY SIMILARITY).
--- STANT BYTHES OF CANCER (BY SIMILARITY).
--- DISEASE: P53 IS FOUND IN SUCHEAR (BY SIMILARITY).
--- STANT BYTHES OF CANCER (BY SIMILARITY).
--- STANT BYTHES OF CANCER (BY SIMILARITY).
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PROSTIE; PS00348; P53; 1.
Activator; Anti-oncogene; Apoptosis; DNA-binding; Nuclear protein; PhosphoryLation; Transcription regulation.
SEQUENCE 387 AA; 43475 MW; 3BB49A432B0053PA CRC64;
                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia! Eutheria; Cetartiodactyla; Suina; Suidae; Sus. NCBI_TaxID=8923;
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                    01-OCT-2000 (TrEMBLrel. 15, Created)
01-OCT-2000 (TrEMBLrel. 15, Last sequence update)
01-UIN-2002 (TrEMBLrel. 21, Last annotation update)
Cellular tumor antigen p53.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Search completed: February 12, 2003, 10:35:38
                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
TISSUE=URINARY BLADDER EPITHELIUM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00870; P53; 1.
PRINTS; PR00386; P53SUPPRESSR.
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Matches 17; Conservative
                                                                                                                                       Sus scrofa (Pig).
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February 12, 2003, 10:49:54; Search time 15 Seconds (without alignments) 32.045 Million cell updates/sec
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                         283224 seqs, 96134422 residues
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Maximum Match 100%
Listing first 100 summaries
                                                          OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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1 FXXLW 5
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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Maximum DB seq length: 5
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Perfect score:
Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

Description	cholecystokinin-5	neuropentide - sea	1.10	bradykinin-potenti	Ig heavy chain CRD	Iq heavy chain CRD	ell re	T-cell receptor be	Iq heavy chain CRD	alkanal monooxygen		andiotensin-conver	ribulose-bisphosph	actin I - malaria		Θ	Met-enkephalin - b	subesophageal gang	cocoonase (EC 3.4.	T-cell receptor be	serrawettin W2 - S	proctolin - Americ	peptidyl-dipeptida	copper resistance	flagellar protein	R-phycoerythrin al		proctolin - Atlant	cal
SUMMARIES	A32516	A60803	JH0253	G37196	PT0281	PT0308	PT0729	PT0580	PT0278	A44955	870615	PQ0009	PS0324	B45525	A44692	B61445	A61445	JS0319	B61168	PT0644	A58728	HOROHA	JN0860	C41225	E42364	B22565	S55237	A60411	853595
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R.Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G. J. Exp. Med. 173, 395-407, 1991
A.Title: Preferential utilization of specific immunoglobulin heavy chain diversity and j A;Reference number: PT0222; MUID:91108337; FMID:1899102
A;Recession: PT0308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PTO291
B;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
J. Exp. Med. 173, 395-407, 1991
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and A;Reference number: PT0222; MUID:91108337; PMID:1899102
                                                                                                                                                                                                                                                                                                                                                                                                                                                   C,Accession: G37156
R,Cintra, A.C.O.; Vieira, C.A.; Giglio, J.R.
T, Forberin Chem. 9, 221-227, 1990
A;Title: Primary structure and biological activity of bradykinin potentiating peptides A;Title: Primary Structure and biological activity of bradykinin potentiating peptides A;Reference number: A37196; MUID:90351557; PMID:2386615
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C;Species: Homo sapiens (man)
C;Date: 30-Sep.1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0308
                                                                                                                                                                                                                                                                                                                                                          bradykinin-potentiating peptide 7 - island jararaca
C;Species: Bothrops insularis (island jararaca)
C;Date: 14-Feb-1992 #sequence_revision 01-Dec-1992 #text_change 05-Aug-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A,Accession: G37196
A,Status: preliminary
A,Molecule type: protein
A,Residues: 1-5 <CIN>
C,Keywords: pyrcoglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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47.8%; Score 11; DB 2; Length 5; ilarity 100.0%; Pred. No. 2.8e+05; Conservative 0; Mismatches 0; Indels
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A, Experimental source: B lymphocyte
C, Keywords: heterotetramer; immunoglobulin
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          Query Match
Best Local Similarity
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A;Experimental source: gut
C;Comment: This peptide increased basal tone of the circular muscle of the esophagogastr , and of the circular muscle of the gastro-intestinal junction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Molecule type: protein
A;Residues: 1-5 <SHI>
C;Comment: This peptide corresponds to the five carboxyl-terminal residues of cholecystd
                                                                                                                                                                                                                                                                                          R;Shively, J.; Reeve Jr., J.R.; Eysselein, V.E.; Ben-Avram, C.; Vigna, S.R.; Walsh, J.H. M. J. Physiol. 252, G272-G275, 1987
A,Title: CCK-5: sequence analysis of a small cholecystokinin from canine brain and intes
A,Reference number: A32516; WUID:87153871; PMID:3826354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Brain Res. 442, 354-358, 1988
A;Title: Isolation of cGiu-Ser-Lu-Arg-Trp-NH-2, a novel neuropeptide from sea anemones.
A;Reference number: A60803; MUID:88222764; PMID:2897223
A;Accession: A60803
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0
                                                                                                                                 cholecystokinin-5 - dog
N;Alternate names: CCK-5
C;Species: Canis lupus familiaris (dog)
C;Date: 18-Oct-1989 #sequence_revision 18-Oct-1989 #text_change 18-Aug-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Species: Anthopleura elegantissima 15pecies: Anthopleura elegantissima 15pecies: Anthopleura elegantissima 15pecies: Ancession: Accession: Accession: Accession: Accession: C.J.P. 67saff, D.; Grimmelikhuijzen, C.J.P.
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A;Residues: 1-5 <GRA>
K;Rewords: amidated carboxyl end; neuropeptide; pyroglutamic acid
F;J/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;S/Modified site: amidated carboxyl end (Trp) #status experimental
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C;Keywords: amidated carboxyl end; neuropeptide
F;5/Modified site: amidated carboxyl end (Phe) #status experimental
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    sea anemone (Anthopleura elegantissima)

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47.8%; Score 11; DB 2; Le
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0;
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47.8%; Score 11; DB 2; Le
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0;
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A; Molecule type: protein
                                                                                                                                                                                                                                                                  Accession: A32516
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A; Residues: 1-5 < YAM>

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PT0278
Ig heavy chain CRD3 region (clone 4-88) - human (fragment)
C;Species: Homo sapiens (man)
C;Decises: Homo sapiens (man)
C;Date: 30-Sep-1993 #sequence_revision 30-Sep-1993 #text_change 16-Aug-1996
C;Accession: PT0278
R;Yamada, M.; Wasserman, R.; Reichard, B.A.; Shane, S.; Caton, A.J.; Rovera, G.
A;Title: Preferential utilization of specific immunoglobulin heavy chain diversity and jakeference number: PT0222; MUID:91108337; PMID:1899102
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      alkanal monooxygenase (FMN-linked) (EC 1.14.14.3) alpha chain - Vibrio harveyi (fragment C;Species: Vibrio harveyi
C;Species: Vibrio harveyi
C;Species: Vibrio harveyi
C;Saccession: A44955
R;Paquatte, O.; Tu, S.C.
R;Paquatte, O.; Tu, S.C.
A;Title: Chemical modification and characterization of the alpha cysteine 106 at the Vibria A;Reference number: A44955; MUID:90175700; PMID:2626493
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NyAlternate names: xylanase
NyAlternate names: xylanase
Species Streptomyces sp.
A;Variety: Chainia sp. NCL 82.5.1
C;Date: 19-Mar-1998 #sequence_revision 17-Apr-1998 #text_change 07-May-1999
C;Accession: S70615
R;Rao, M.; Khadilkar, S.; Bandivadekar, K.R.; Deshpande, V.
Biochem. J. 316, 771-775, 1996
A;Title: Structural environment of an essential cysteine residue of xylanase from Chainia, A;Reference number: S70615; MUID:96265041; PMID:8670151
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A;Description: endohydrolyzation of beta-1,4-xylosidic linkages in xylans
A;Pathway: fermentation of hemicellulose into ethanol
C;Keywords: glycosidase; hydrolase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 8; DB 2; Length 5;
Pred. No. 2.8e+05;
0; Mismatches 2; Indels
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A,Molecule type: protein
A,Residues: 1-5 <PAQ>
C,Keywords: FMN; luminescence; monooxygenase; oxidoreductase
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A,Molecule type: protein
A,Residues: 1-5 cRAO.
A,Experimental source: Chainia sp. strain NCL 82.5.1
A,Note: the source is designated as Chainia sp.
                                                                                                                                                                                                                                                                                                                                                        A;Molecule type: DNA
A;Residues: 1-5 <PAM>
A;Experimental source: B lymphocyte
C;Keywords: heterotetramer; immunoglobulin
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50.0%;
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Best Local Similarity 50.0
Matches 2; Conservative
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A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reterence number: PT0509; MUID:91277601; PMID:1711558
A;Recession: PT0640
A;Status: translation not shown
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C;Species: Mus musculus (house mouse)
C;Decies: Mus musculus (house mouse)
C;Decession: PT0589
R;Feeney, A.J.
J. Exp. Med. 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Species: Was musculus (house mouse)
C;Date: 17-Jul.1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0640; PT0685; PT0729
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A/Residues: 1-5 <FES.
A/Experimental source: newborn thymus, strain BALB/c, clone 120-1J
A/Accession: PT0685
A/Status: translation not shown
A/Molecule type: DNA
A/Residues: 1-5 < FE2.
A/Experimental source: day 18 fetal thymus, strain BALB/c, clone 154-1C
A/Accession: PT0729
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A;Molecule type: DNA
A;Residues: 1-5 <FE3.
A;Experimental source: newborn thymus, strain BALB/c, clone 135-1AG
C;Keywords: T-cell receptor
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A;Residues: 1-5 <FEE>
A;Experimental source: day 19 fetal thymus, strain BALB/c
C;Keywords: T-cell receptor
A; Experimental source: B lymphocyte C; Keywords: heterotetramer; immunoglobulin
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Matches 1; Conservative
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C;Species: Achatina fulica (giant African snail)
C;Date: 23-Mar-1995 #sequence_revision 05-Apr-1995 #text_change 11-Jul-1997
C;Date: 23-Mar-1995 #sequence_revision 05-Apr-1995 #text_change 11-Jul-1997
C;Accession: 444692
C;Accession: A44692
C;Accession: A44692
C;Accession: A44692
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C;Species: Mytilus edulis (blue mussel)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 21-Jan-2000
C;Accession: B61445
R;Leung, M.K.; Stefano, G.B.
R;Leung, M.K.; Stefano, G.B.
R;Leung, M.K.; Stefano, G.B.
A;Title: Isolation and identification of enkephalins in pedal ganglia of Mytilus edulis
A;Reference number: A61445; MUID:84144823; PMID:6583690
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Met-enkephalin - blue mussel
Species: Mytilus edulis (blue mussel)
C;Species: Mytilus edulis (blue mussel)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 21-Jan-2000
C;Accession: A61445
Escfano, G.B.
Proc. Natl. Acad. Sci. U.S.A. 81, 955-958, 1984
A;Title: Isolation and identification of enkephalins in pedal ganglia of Mytilus edulis
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A;Residues: 1-5 <OHT>
C;Keywords: amidated carboxyl end; D-amino acid; neuropeptide
F;Z/Modified site: D-asparagine (Asn) #status experimental
F;S/Modified site: amidated carboxyl end (Val) #status experimental
                      A; Note: the authors translated the codon GAA for residue 3 as Gly C; {\tt Comment}: The actin I gene contains no introns.
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                                                                                                                 Ouery Match 26.1%; Score 6; DB 2; Length 5; Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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A;Residues: 1-5 <LEU>
A;Experimental source: pedal ganglia
C;Keywords: neuropeptide; opioid peptide
A;Cross-references: GB:J03988
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C;Species: Plasmodium falciparum
C;Species: Plasmodium falciparum
C;Species: Plasmodium falciparum
C;Species: Plasmodium falciparum
C;Species: O3-Unr.1993 #sequence_revision 28-Oct-1994 #text_change 09-Jun-2000
C;Accession: B45525
R;Wesseling, J.G.; Snijders, P.J.F.; van Someren, P.; Jansen, J.; Smits, M.A.; Schoenmak
Mol. Biochem. Parasitol. 35, 167-176, 1989
A;Titles: Stage-specific expression and genomic organization of the actin genes of the ma
A;Accession: B45525; MUID:89364996; PMID:2671721
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R; Maruyama, S.; Miyoshi, S.; Tanaka, H.
Agric. Biol. Chem. 53, 2763-2767, 1989
A; Title: Angiotensin I-converting enzyme inhibitors derived from Ficus carica.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'Species: Oryza_sativa (rice)
.Date: 03-Feb-1994 #sequence_revision 03-Feb-1994 #text_change 02-Jul-1998
                                                                                                                                                                                                                                                                                                                                                               angiotensin-converting enzyme inhibitor (FLP-2) - common fig
NiAlternate names: figus latex peptide 2
C;Species: Ficus carica (common fig)
C;Species: 07-Jun-1990 #sequence_revision 07-Jun-1990 #text_change 08-Dec-1995
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                             Query Match 26.1%; Score 6; DB 2; Length 5; Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 1; Conservative 0; Mismatches 0; Indele
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A;Residues: 1-5 <TSU>
A;Experimental source: leaf, chlorophyll
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A, Residues: 1-5 <MAR>
A, Experimental source: latex
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A;Molecule type: DNA
A;Residues: 1-5 <WES>
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Comp. Biochem. Physiol. C 88, 185-187, 1987
A;Title: Isolation and structure of a peptide isolated from the suboesophageal ganglion
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A:Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
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C;Species: Antheraea pernyi (Chinese oak silkmoth)
C;Species: Antheraea pernyi (Chinese oak silkmoth)
C;Accession: B61168
R;Kramer, K.J.; Felsted, R.L.; Law, J.H.
J. Biol. Chem. 248, 3021-3028, 1973
A;Fitle: Cocoonase. V. Structural studies on an insect serine protease.
A;Feference number: A61168; MUID:73166540; PMID:4735570
                                                                                                                                                                                                                                                                                                                                                                                    subesophageal ganglion pentapeptide - house cricket
C;Species: Acheta domesticus (house cricket)
C;Date: 07-Sep.1990 #sequence_revision 07-Sep-1990 #text_change 20-Jun-2000
C;Accession: JS0319
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
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A;Residues: 1-5 <KRA>
C;Keywords: hydrolase; serine proteinase; zymogen
F;1-5/Product: cocoonase (fragment) #status experimental <MAT>
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A; Reference number: A61445; MUID:84144823; PMID:6583690 A; Accession: A61445
                                     A;Molecule type: protein
A;Residues: 1-5 <LEU>
A;Experimental source: pedal ganglia
C;Keywords: neuropeptide; opioid peptide
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Best Local Similarity
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R; Feeney, A.J.
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C;Species: Serratia marcescens
C;Date: 10-Feb-1998 #sequence_revision 12-Feb-1998 #text_change 12-Feb-1998
C;Accession: A58728
R;Matsuyama, T.; Kaneda, K.; Nakagawa, Y.; Isa, K.; Hara-Hotta, H.; Yano, I.
Bacteriol. 174, 1769-1776, 1992
A;Title: A novel extracellular cyclic lipopeptide which promotes flagellum-dependent and A;Reference number: A58728; MUID:92193260; PMID:1548227
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A;Residues: 1-5 < MAT>
C;Commental source: strain NS 25
C;Comment: A surfactant lipopeptide promoting flagellum-independent surface translocation C;Ceywords: blocked amino end; blocked carboxyl end; D-amino acid; lipoprotein; unencode F;I/Modified site: D-leucine (Leu) #stratus experimental
F;I/Modified site: D-phenylalamine (Phe) #status experimental
F;I/Modified site: D-phenylalamine end, ester carboxyl end (Leu-Ile) #status experimental
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C;Species: Periplaneta americana (American cockroach)
C;Date: 29-Jul-1981 #sequence_revision 29-Jul-1981 #text_change 23-Aug-1996
C;Accession: A01644
R;Starratt, A.N.; Brown, B.E.
Life Sci. 17, 1253-1256, 1975
A;Title: Structure of the pentapeptide proctolin, a proposed neurotransmitter in insects A;Reference number: A93048; MUID:76074708; PMID:576
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A,Residues: 1-5 <STA>
A,Note: the synthetic peptide had the same chromatographic, electrophoretic, and pharmac
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C; Comment: This peptide is found in the lateral white neurons, which occur (in the cockra
innervate the striated hindgut muscles in insects and stimulate contraction of these mus
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Science 213, 567-569, 1981
A;Title: Pentapeptide (proctolin) associated with an identified neuron.
A;Reference number: A94260; MUID:81225865; PMID:6113690
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A; Reference number: PT0509; MUID:91277601; PMID:1711558
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Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0;
                                A;Accession: PT0644
A;Status: translation not shown
A;Molecule type: mRNA
A;Residues: 1-5 <FEE>
A;Experimental source: newborn thymus, strain BALB/c
C;Keywords: T-cell receptor
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100.0%; Pred. No. 2.8
tive 0; Mismatches
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Best Local Similarity
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zinc_binding protein ZBP14 - maize (fragment)

zinc_binding protein ZBP14 - maize (fragment)

C;Species: Zea mays (maize)

C;Species: Zea mays (maize)

C;Date: 2.7-oct-1995 #sequence_revision 03-Nov-1995 #text_change 07-May-1999

C;Accession: S55237

R;Robinson, K.; Jones, D.; Howell, S.; Soneji, Y.; Martin, S.; Aitken, A.

B;Chem. J. 307, 267-272, 1995

A;Reference number: S55237; MUID:95234046; PMID:7717986

A;Accession: S55237

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R;Groome, J.R.; Tillinghast, B.K.; Townley, M.A.; Vetrovs, A.; Watson III, W.H.; Hunt, D Beptides 11, 205-211, 1990
A;Title: Identification of proctolin in the central nervous system of the horseshoe crab A;Reference number: A60411; MUID:90287800; PMID:2356151
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A;Residues: 1-5 <05RO>
C;Comment: This neuropeptide stimulates cardiac output and hindgut motility in the horse
C;Keywords: neuropeptide
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C;Species: Limulus polyphemus (Atlantic horseshoe crab)
C;Date: 03-Feb-1993 #sequence_revision 03-Feb-1993 #text_change 17-Mar-1999
                                                                                                                                                                                                                                                                                                                                             C;Species: Gastroclonium coulteri
C;Date: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993
C;Accession: B22565
R;Klotz, A.V.; Glazer, A.N.
J Biol. Chem. 260, 4856-4863, 1985
A;Tile: Chem. 260, 4856-4863, 1985
A;Reference number: A22565; MUID:85182601; PMID:3886644
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                      Length 5;
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                   17.4%; Score 4; DB 2; Ler
llarity 100.0%; Pred. No. 2.8e+05;
Conservative 0; Mismatches 0;
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Best Local Similarity
Matches 1; Conservat
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A;Residues: 1-5 <KLO>
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R, Matsumura, N.; Fujii, M.; Takeda, Y.; Shimizu, T.
Biosci. Blockchnol. Biochem. 57, 1743-1744, 1993
B. Stille: Isolation and characterization of angiotensin I-converting enzyme inhibitory pe A; Reference number: JN0859; MUID: 94080036; PMID: 7764272
A; Reference number: JN0860
A; Molecule type: protein
A; Residues: 1-5 < MAT.
A; Residues: 1-5 < MAT.
A; Residues: The carboxyl-terminus is essential for the protein's expression of angiotensic C; Superfamily: bradykinin-potentiating peptide
C; Superfamily: bradykinin-potentiating peptide
C; Keywords: angiotensin-converting enzyme inhibitor
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R;Cha, J.S.; Cooksey, D.A.
R;Cha, J.S.; Cooksey, D.A.
Aproc. Natl. Acad. Sci. U.S.A. 88, 8915-8919, 1991
A;Title: Copper resistance in Pseudomonas syringae mediated by periplasmic and outer mem A;Reference number: A41225; MUID:92020961; PMID:1924351
A;Accession: C41225
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J. Bacteriol. 173, 3564-3572, 1991
A;Fitle: Salmonella typhimurium mutants defective in flagellar filament regrowth and A;Reference number: A42364; MUID:91258342; PMID:1646201
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C.Date: 10-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 07-May-1999
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C.Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 24-Jun-1993
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C;Date: 24-Jul-1992 #sequence_revision 24-Jul-1992 #text_change 30-Sep-1993
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  Gaps
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  Indels
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0; Mismatches
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Best Local Similarity
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A;Residues: 1-5 <VOG>
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  Matches
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A; Status: translation not shown
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                                                                                                    Query Match
Best Local Similarity 1
         A; Molecule type: mRNA
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G44817
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                                                                                                                                                                                                                              hypothetical protein (upstream of transcription factor, CCAAT-binding) - chicken C;Species: Gallus gallus (chicken)
C;Species: Gallus gallus (chicken)
C;Date: 15-Jul-1995 #sequence_revision 01-Sep-1995 #text_change 07-May-1999
C;Accession: S53595
R;Calkhoven, C.F.; Bouwman, P.R.J.; Snippe, L.; Ab, G.
Nucleic Acids Res. 22, 5540-5547, 1994
A;Title: Translation start site multiplicity of the CCAAT/enhancer binding protein alpha A;Reference number: S53595; MUID:95140613; PMID:7838705
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NyAlternate names: superprotein peptide 2

C;Species: Oryctolagus cuniculus (domestic rabbit)

C;Date: 31-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 15-Jun-1996

C;Date: 31-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 15-Jun-1996

B;Locewy, A.; Liu, W.S.; Battinger, C.; Willard, M.B.

J. Neurosci. 11, 3412-3421, 1991

A;Reference number: A44823; MUID:92044785; PMID:1941090

A;Accession: D44823
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0624
A;Ribeney, A.J.
J. Exp. Med. 174, 115-124, 1991
A;Riberence number: PT0509; MUID:91277601; PMID:1711558
A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Retus: translation not shown
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                             Indels
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17.4%; Score 4; DB 2; Length 5;
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0; Indel
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A;Molecule type: protein
A;Residues: 1-5 <LOE;
A;Reperimental source: visual tissue
A;Reperimental source: visual tissue
A;Note: sequence extracted from NCBI backbone (NCBIP:64250)
C;Keywords: membrane trafficking
                             0,
Best Local Similarity 100.0%; Pred. No. 2.8e+05; Matches 1; Conservative 0; Mismatches 0;
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A;Molecule type: DNA
A;Residues: 1-5 <CAL>
A;Cross-references: BMBL:X66844
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C;Accession: PT0625 Frogram of the property A.J.
R;Feeney, A.J.
J. Exp. Med. 114, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reference number: PT0509; MUID:91277601; PMID:1711558
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A;Title: Unictional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
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C;Species: Leuconostoc oenos phage P32
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C;Accession: G44817
R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0713
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100.0%; Pred. No. 2.8e+05;
tive 0; Mismatches 0; Indels
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17.4%; Score 4; DB 2; Length 5;
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0; Indel.
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17.4%; Score 4; DB 2; Ler
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0;
A;Residues: 1-5 <FEE>
A;Experimental source: newborn thymus, strain BALB/c
C;Keywords: T-cell receptor
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A;Molecule type: DNA
A;Residues: 1-5 cFEE.
A;Experimental source: newborn thymus, strain BALB/C
C;Keywords: T-cell receptor
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A;Experimental source: newborn thymus, strain BALB/c
C;Keywords: T-cell receptor
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RESULT 40
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C,Species: Leuconostoc oenos phage P37
C,Date: 31-Max-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C,Accession: 144817
R,Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A,Richerence number: A44817; MUID:92085033; PMID:1748868
A,Accession: 144817
A,Molecule type: protein
A,Molecule type: protein
A,Residues: 1-5 <ARE>
A,Residues: 1-5 <ARE>
A,Note: sequence extracted from NCBI backbone (NCBIP:70330)
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C;Species: Leuconostoc oenos phage P54
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C;Accession: E44817
R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J, Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
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C;Species; Leuconostoc oenos phage PAt5-12
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
C;Accesion: C44817. Lonvaud, A.; Hammes, W.P.
B;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
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17.4%; Score 4; DB 2; Length 5;
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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17.4%; Score 4; DB 2; Length 5;
Best Local Similarity 100.0%; Pred. No. 2.8e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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J. Gen. Microbiol. 137, 2135-2139, 1991
A; Title: Lysogeny in Leuconostoc oenos.
A; Reference number: A44817; MUID:92085033; PMID:1748868
A; Accession: G44817
A; Molecule type: protein
A; Residues: 1-5 < ARE>
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A;Residues: 1-5 <ARE>
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C;Species: Leuconostoc oenos phage PAt5-12
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
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C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
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C;Species: Leuconostoc oenos phage PZtl1-15
C;Date: 31-Mar-1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
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A,Residues: 1-5 <ARE>
A,Note: sequence extracted from NCBI backbone (NCBIP:70342)
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: G44817
A;Molecule type: protein
A;Residues: 1-5 < ARE>
A;Note: sequence extracted from NCBI backbone (NCBIP:70341)
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R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Tille: Lysogeny in Leuconostoc Oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
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R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: B44817
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A;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J; Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
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ibosomal protein S4 - Bacillus circulans (fragment)
C;Species: Bacillus circulans
C;Species: 19-01-1996 #sequence_revision 19-01-1996 #text_change 19-01-1996
C;Date: 19-01-13996 #sequence_revision 19-01-1996 #text_change 19-01-1996
C;Accession: 133964 #sequence_revision 19-01-1996 #text_change 19-01-1996
C;Accession: 133964 #sequence_revision of the Bacillus subtilis rpsD regulatory target site.
A;Reference number: 139963; MUID:93015735; PMID:1400226
A;Accession: 139964
A;Accession: 139964
A;Accession: 159964
A;Residues: 1-5 cRES>
A;Residues: 1-5 cRES>
A;Construction of the Bacillus subtilis rpsD regulatory target site.
A;Residues: 1-5 cRES>
A;Construction of the Bacillus subtilis rpsD regulatory target site.
A;Residues: 1-5 cRES>
A;Construction of the Bacillus subtilis rpsD regulatory target site.
A;Residues: 1-5 cRES>
A;Construction of the Bacillus subtilis rpsD regulatory target site.
A;Residues: 1-5 cRES>
A;Construction of the Bacillus subtilis rpsD regulatory target site.
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C;Species: Bacillus licheniformis
C;Species: 19-Jul-1996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1996
C;Accession: 13996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1996
R;Grundy, F.J.; Henkin, T.M.
J. Bacteriol. 174, 6763-6770, 1992
A;Title: Characterization of the Bacillus subtilis rpsD regulatory target site.
A;Reference number: 139963; MUID:93015735; PMID:1400226
A;Accession: 139966
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13.0%; Score 3; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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A;Residues: 1-5 <RES>
A;Cross-references: GB:M99043; NID:g143475
C;Genetics:
                                                     A;Status: preliminary
A;Molecule type: DNA
A;Residues: 1-5 <BRA>
A;Cross-references: GB:M63747
                    A; Accession: B37325
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C;Species: Staphylothermus marinus
C;Date: 04-Dec-1997 #sequence_revision 04-Dec-1997 #text_change 17-Mar-1999
C;Date: 04-Dec-1997 #sequence_revision 04-Dec-1997 #text_change 17-Mar-1999
C;Accession: S69237
Nol. Biol. 245, 385-401, 1995
J. Mol. 2
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C;Date: 11-Sep-1992 #sequence_revision 11-Sep-1992 #text_change 23-Mar-1993
C;Accession: B37325
R;Braaten, B.A.; Blyn, L.B.; Skinner, B.S.; Low, D.A.
J. Bacteriol. 173, 1789-1800, 1991
A;Title: Evidence for a methylation-blocking factor (mbf) locus involved in pap pilus ex
A;Reference number: A37325; MUID:91154136; PMID:1671857
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Proc. Natl. Acad. Sci. U.S.A. 93, 7623-7627, 1996
A;Title: Phytosulfokine, sulfated peptides that induce the proliferation of single mesop
A;Reference number: JT0870
A;Accession: JT0870
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C;Species: Asparagus officinalis (garden asparagus)
C;Date: 02-Apr-2002 #sequence_revision 02-Apr-2002 #text_change 19-Apr-2002
C;Accession: JT0870
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C;Species: Escherichia coli
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                                                                     A;Molecule type: protein
A;Residues: 1-5 <ARE>
A;Note: sequence extracted from NCBI backbone (NCBIP:70340)
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A;Residues: 1-5 <PET>
A;Experimental source: strain F1, DSM 3639
C;Keywords: cell wall; glycoprotein; heat-stable protein
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: D44817
                                                                                                                                                                                                                                                                                                               1; Conservative
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NyAlernate names: Abcp-B
C;Species: Ascidia ceratodes
C;Date: 19-Mar-1997 #sequence_revision 29-Aug-1997 #text_change 11-Jun-1999
C;Accession: 868326
R;Taylor, S.W.; Ross, M.M.; Waite, J.H.
Rxch. Biochem. Biophys. 324, 228-240, 1995
A;Title: Novel 3,4-di- and 3,4,5-trihydroxyphenylalanine-containing polypeptides from th
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C;Species: 19-May-1994 #sequence_revision 19-May-1994 #text_change 17-Mar-1999 C;Species: 19-May-1994 #sequence_revision 19-May-1994 #text_change 17-Mar-1999 C;Accession: PQ0689 F; Dyboxata, J; Mikami, K; Hayashida, N; Nakamura, M.; Sugiura, M. Plant Physiol. 102, 1259-1267, 1993 A;Title: Molecular heterogeneity of photosystem I. psab, psab, psab, psab, psab, are A;Reference number: PQ0667; MUID:94105345; PMID:8278548
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A;Residues: 1-5 <17A?
-2,Modified site: 3',4',5'-trihydroxyphenylalanine (Tyr) #status experimental
F;4/Modified site: 3',4'-dihydroxyphenylalanine (Tyr) #status experimental
                                                                                     P.phycoerythrin gamma-A chain - red alga (Gastroclonium coulteri) (fragment) (Species: Gastroclonium coulteri C,Species: Gastroclonium coulteri C,Species: Gastroclonium coulteri C,Species: Gastroclonium coulteri C,Species: Anar-1998 #sequence_revision 07-Mar-1998 #text_change 23-Mar-1993 C,Accession: F22565 A.N. ; Glazer, A.N. ; Glazer, A.N. ; Glazer, A.N. ; Glazer, A.N. ; A.N. ; Glazer, A.N. ; A.Y. ; Licle: Characterization of the blinn attachment sites in R-phycoerythrin. A,Reference number: A22565, MUID:85182601; PMID:3886644 A,Molecule type: protein A,Nolecule type: protein A,Residues: 1-5 <KLO>
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A;Residues: 1-5 <080>
C;Keywords: chloroplast; photosynthesis; photosystem I; thylakoid
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Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0
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R;Nagai, S.; Wiker, H.G.; Harboe, M.; Kinomoto, M.
Infect. Immun. 59, 372-382, 1991
A;Title: Isolation and partial characterization of major protein antigens in the culture A;Reference number: A60274; MUID:91099989; PMID:1898899
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             dnaZX-like protein - Bacillus subtilis (fragment)
C;Species: Bacillus subtilis
C;Species: Bacillus subtilis
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 21-Jul-2000
C;Accession: 140469
R;Struck, J.C.; Hartmann, R.K.; Toschka, H.Y.; Erdmann, V.A.
Mol. Gen. Genet. 215, 478-482, 1989
A;Title: Transcription and processing of Bacillus subtilis small cytoplasmic RNA.
A;Reference number: 140469; MUID:89218958; PMID:2468993
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                                C;Accession: I39965
R;Grundy, F.J.; Henkin, T.M.
J. Bacteriol. 174, 6763-6770, 1992
A;Reference number: 139963; MUID:93015735; PMID:140026
A;Reference number: 139963; MUID:93015735; PMID:1400226
A;Accession: 139963; MUID:93015735; PMID:140026
A;Accession: 139963
A;Molecule type: DNA
A;Residues: 1-5 <RES>
A;Cenetics: 1-5 <RES>
A;Genetics: 1-5 <RES>
A;Genetics: A;Ge
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C;Species: Mycobacterium tuberculosis
C;Date: 11-Dec-1992 #sequence_revision 11-Dec-1992 #text_change 30-Sep-1993
C;Date: 19-Jul-1996 #sequence_revision 19-Jul-1996 #text_change 19-Jul-1996
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A;Molecule type: DNA
A;Residues: 1-5 <RES>
A;Cross-references: EMBL:X14796; NID:g40130; PIDN:CAA32902.1; PID:g4376204
C;Genetics:
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primase - Citrobacter diversus (fragment)
C;Species: Citrobacter diversus
C;Species: Citrobacter diversus
C;Species: Citrobacter diversus
C;Species: Citrobacter diversus
C;Species: 140702
R;Versalovic, J.; Lupski, J.R.
Mol. Microbiol. 8, 343-355, 1993
Mol. Microbiol. 8, 343-355, 1993
A;Title: Conservation and evolution of the rpsU-dnaG-rpoD macromolecular synthesis (MMS)
A;Reference number: 140702; MUID:93302510; PMID:8316085
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                                                                                                                                                                                                                  C;Species: Xanthomonas sp. C;Date: 15-Feb-1997 #text_change 08-Oct-1999 C;Date: 15-Feb-1997 #sequence_revision 13-Mar-1997 #text_change 08-Oct-1999 C;Accession: 870154 #sequence_revision 13-Mar-1997 #text_change 08-Oct-1999 R;Kholodii, G.Y.; Mindlin, S.Z.; Bass, I.A.; Yurieva, O.V.; Minakhina, S.V.; Nikiforov, Mol. Microbiol. 17, 1189-1200, 1995 A;Title: Four genes, two ends, and a res region are involved in transposition of Th5053: A;Reference number: 870140; MUID:96130850; PMID:8594337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: DNA
A;Residues: 1-5 < KHO>
A;Cross-references: EMBL:L40585; NID:g710572; PIDN:AAA98329.1; PID:g735909
A;Note: the nucleotide sequence was submitted to the EMBL Data Library, May 1995
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8.7%; Score 2; DB 2; L
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0
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A;Molecule type: DNA
A;Residues: 1-5 <RES>
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                                                                                                                                                                                           URF2 protein - Xanthomonas sp.
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Matches
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                    glycogen phosphorylase (EC 2.4.1.1), muscle - mullet (Liza ramada) (fragment)
N.Alternate names: glycogen phosphorylase b
C;Species: Liza ramada
C;Species: Liza ramada
C;Species: Liza ramada
C;Accession: A60521
R;Bonamusa, L.; Baanante, I.V.
Comp. Biochem. Physiol. B 95, 295-301, 1990
A;Title: Purification and characterization of glycogen phosphorylase B from skeletal muscle paracterization: A60521
A;Accession: A60521
A;Accession: A60521
A;Accession: A60521
                                                                                                                                                                                                                                                                                                                      A,Molecule type: protein
A,Residues: 1-5 <BON>
C,Superfamily: phosphorylase
C,Keywords: glycosyltransferase, hexosyltransferase, phosphoprotein
F,3/Binding site: phosphate (Ser) (covalent) (by phosphorylase b kinase) #status experin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       G;Species: Sarda orientalis (striped bonito)
C;Date: IO-Mar-1994 #sequence_revision 10-Mar-1994 #text_change 07-May-1999
C;Accession: JN0862
R;Matsumura, N.; Fujii, M.; Takeda, Y.; Shimizu, T.
Bibsci: Bibtechnol, Biochem. S7, 1743-11744, 1993
A;Title: Isolation and characterization of angiotensin I-converting enzyme inhibitory peareference number: JN0859; MUID:94080036; PMID:7764272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A) Experimental source: intestine C; Comment: The amino terminal tripeptide of this protein inhibits angiotensin I-converti C; Superfamily: bradykinin-potentiating peptide C; Keywords: angiotensin-converting enzyme inhibitor
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C;Species: Pseudomonas syringae pv. tomato
C;Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 24-Jun-1993
C;Accession: B41225
R;Cha, J.S.; Cooksey, D.A.
Proc. Natl. Acad. Sci. U.S.A. 88, 8915-8919, 1991
A;Title: Copper resistance in Pseudomonas syringae mediated by periplasmic and outer men A;Reference number: A41225; MUID:92020961; PMID:1924351
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0.0%; Pred. No. 2.8e+05;
7ative 1; Mismatches 0; Indels
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0.0%; Pred. No. 2.8e+05;
rative 1; Mismatches 0; Indels
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0.0%; Pred. No. 2.8e+05;
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Matches 0; Conservative
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A;Molecule type: protein
A;Residues: 1-5 <MAT>
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A;Molecule type: protein
A;Residues: 1-5 <CHA>
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C;Date: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 21-Jul-2000 C;Accession: T14910 Fsequence_revision 20-Sep-1999 #text_change 21-Jul-2000 C;Accession: T14910 Fsequence_revision 20-Sep-1999 Fskircher, S.; Ledger, S.; Hadger, H.; Weisshaar, B.; Schafer, E.; Frohnmeyer, H. Mol. Gen. Genet. 257, 595-605, 1998 A;Title: CPRF4a, a novel plant bZIP protein of the CPRF family: comparative analysis of A;Reference number: 218261; MUID:98265918; PMID:9604882
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   C;Accession: B37988
R;Murakami-Murofushi, K.; Takahashi, T.; Minowa, Y.; Iino, S.; Takeuchi, T.; Kitagaki-Og
N; Biol. Chem. 265, 19898-19903, 1990
A;Title: Purification and characterization of a novel intracellular acid proteinase from A;Reference number: A37988; MUID:91060608; PMID:2246266
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C'Date: 28-Mar-1991 #sequence_revision 28-Mar-1991 #text_change 23-Jun-1993
C'Accession: A37114
R'Yuan, L.; Craig, S.P.; McKerrow, J.H.; Wang, C.C.
A' Blol. Chem. 265, 13528-13532, 1990
A'Title: The hypoxanthine phosphoribosyltransferase of Schistosoma mansoni. Furt
A;Reference number: A37114; MUID:90337955; PMID:2199439
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                          A;Cross-references: EMBL:Y10809; NID:g3336901; PIDN:CAA71767.1; PID:g3336902 A;Experimental source: Hamburger Schnitt
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A,Cross-references: EMBL:Y10810; NID:g3336904; PIDN:CAA71769.1; PID:g3336905
A,Experimental source: ssp. Hamburger Schnitt
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8.7%; Score 2; DB 2; Length 5;

Best Local Similarity 0.0%; Pred. No. 2.8e+05;

Matches 0; Conservative 1; Mismatches 0; Indels
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8.7%; Score 2; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               hypothetical protein - parsley
C;Species: Petroselinum crispum (parsley)
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A; Residues: 1-5 <KIR>
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C;Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 28-May-1999
C;Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 28-May-1999
C;Date: 31-Mar-1990 #sequence_revision 31-Mar-1990 #text_change 28-May-1999
R;Anderson, B.E.; Baumstark, B.R.; Bellini, W.J.
J; Bacteriol. 170, 4493-4500, 1988
A;Title: Expression of the gene encoding the 17-kilodalton antigen from Rickettsia ricke
A;Reference number: A91885; MUID:89008059; PMID:3139629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          C;Accession: T10954
R;Christiansen, A.; Hansen, A.C.; Vijn, I.; Pallisgard, N.; Larsen, K.; Yang, W.C.; Biss submitted to the EMBL Data Library, December 1995
A;Description: A novel type of DNA binding protein interacts with a conserved sequence in A;Reference number: 217228
A;Accession: T10954
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('Species: Petroselinum crispum (parsley)
('Species: Petroselinum crispum (parsley)
('Species: Petroselinum crispum (parsley)
('Species: 20-Sep-1999 #sequence_revision 20-Sep-1999 #text_change 21-Jul-2000
('Accession: T14908
R'Kircher, S.; Ledger, S.; Hayashi, H.; Weisshaar, B.; Schafer, E.; Frohnmeyer, H.
Mol. Gen. Genet. 257, 595-605, 1998
A;Title: (CRR#4a, a novel plant bZIP protein of the CPRF family: comparative analysis of A;Reference number: Z18261; MUID: 98265918; PMID: 9604882
A;Accession: T14908
A;Accession: T14908
A;Status: preliminary; translated from GB/EMBL/DDBJ
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C;Species: Vicia sativa (spring vetch, tare)
C;Date: 16-Jul-1999 #sequence_revision 16-Jul-1999 #text_change 16-Jul-1999
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                                                                                                                                   0;
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                                                                         Length 5;
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                                                                   Parch 8.7%; Score 2; DB 2; Local Similarity 0.0%; Pred. No. 2.8e+05; es 0; Conservative 1; Mismatches 0
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      A; Cross-references: GB: M21922; NID: g144434
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C;Accession: B44823
R;Loewy, A; Liu, W.S.; Baitinger, C.; Willard, M.B.
N. Neurososci. 11, 3422-3421, 1991
A;Title: The major 35S-methionine-labeled rapidly transported protein (superprotein) is A;Reference number: A44823; MUID:92044785; PMID:1941090
A;Reference number: A44823; MUID:92044785; PMID:1941090
A;Reference number: A44823
A;Status: precliminary
A;Residues: 15 < LOE>
A;Residues: 15 < 
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R;Fencey, A.J.
Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          J. Exp. Med. 174, 115-124, 1991
A;Title: Unretional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Accession: PT0597
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
C;Accession: PT0597
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C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
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8.7%; Score 2; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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8.7%; Score 2; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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C;Keywords: T-cell receptor
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A;Residues: 1-5 cFEB.
A;Experimental source: newborn thymus, strain BALB/c
C;Keywords: T-cell receptor
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A;Molecule type: mRNA
A;Residues: 1-5 <FEE>
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(Species: Gallus gallus (chicken)

(Jobate: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 28-Feb-1997

(Jobate: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 28-Feb-1997

(Jobate: 13-Sep-1996 #sequence_revision 13-Sep-1996 #text_change 28-Feb-1997

(Jobate: 15-Sep-1996 #sequence_revision 150385

R;Shen, R.; Goswami, S.K.; Mascareno, E.; Kumar, A.; Siddiqui, M.A.Q.

R;Shen, R.; Goswami, S.K.; Mascareno, E.; Kumar, A.; Siddiqui, M.A.Q.

A;Title: Tissue-specific transcription of the cardiac myosin light-chain 2 gene is regul A;Reference number: 150385

A;Accession: 150385

A;Accession: 150385

A;Accession: preliminary; translated from GB/EMBL/DDBJ

A;Residues: 1-5 <SHE>
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FBBS Lett. 382, 15-17, 1996
A;Title: Crystallization and preliminary X-ray diffraction analysis of boar seminal plas A;Reference number: 862882; MUID:96196555; PMID:8612739
A;Accession: 862883
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A;Residues: 1-5 <ROM>
C;Complex: heterodimer; seminal plasma protein I and seminal plasma protein
C;Keywords: glycoprotein; heterodimer; semen
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N,Alfernate names: superprotein peptide 10A
C,Species: Oryctolagus cuniculus (domestic rabbit)
C,Species: Oryctolagus cuniculus (domestic rabbit)
C,Date: 31-Mar-1993 #sequence_revision 18-Nov-1994 #text_change 15-Jun-1996
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C;Species: Sus scrofa domestica (domestic pig)
C;Date: 28-Oct-1996 #sequence_revision 13-Mar-1997 #text_change 17-Mar-1999
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                                       A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-5 <YUA>
C;Keywords: glycosyltransferase; pentosyltransferase
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Best Local Similarity 0.0%
Matches 0; Conservative
           A; Accession: A37114
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I50385
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phosphoprotein, bone - chicken (fragment)
C;Species: Gallus gallus (chicken)
C;Dete: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 04-Mar-2000
C;Date: 21-Nov-1993 #sequence_revision 10-Nov-1995 #text_change 04-Mar-2000
C;Accession: S11127; S11128
R;Mikuni-Takagaki, Y.; Glimcher, M.J.
Biochem. J. 268, 588-591, 1990
A;Title: Post-translational processing of chicken bone phosphoproteins. Identification A;Reference number: S11127; MUID:90303246; PMID:2363696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        spinal cord peptide SCP-6 - pig
Cispecies: Sus scroffa domestica (domestic pig)
Cibate: 28-89-1987 #sequence_revision 28-8ep-1987 #text_change 18-Aug-2000
CiAccession: C23751
Rihsi, K.L.; Chen, R.L.; Chen, Z.G.; Zhang, H.L.; Lu, Y.A.; Guo, S.Y.; Wu, S.X.; Tsou,
Arch. Biochem. Biophys. 240, 178-183, 1985
A;Reference number: A23751; MUID:85250425; PMID:4015098
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C;Species: Mus musculus (house mouse)
C;Addes: 19-Nov-1988 #sequence_revision 18-Aug-2000 #text_change 18-Aug-2000
C;Adcession: A26830
R;Reichelt, K.; Elgjo, K.; Edminson, P.D.
R;Reichelt, K.; Elgjo, K.; Edminson, P.D.
A;Title: Biophys. Res. Commun. 146, 1493-1501, 1987
A;Title: Isolation and structure of an epidermal mitosis inhibiting pentapeptide.
A;Reference number: A26830; MUID:87298602; PMID:3619940
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4.3%; Score 1; DB 2; L
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0
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A;Molecule type: protein
A;Residues: 'X', 2-5 <MIK2>
C;Keywords: phosphoprotein
                                                                                                                                                                                                                                                                                                                                            A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-5 <MIKL>
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A;Molecule type: protein
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                                                                                                                                                                                          J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions.
A;Reference number: PT0509; MUID:91277601; PMID:1711558
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C;Adres: Leuconosico cenos phage P32
C;Adres: Jahar-1934 #sequence_revision 22-May-1998 #text_change 22-May-1998
C;Adcession: H44817
B;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostco cenos
A;Title: Lysogeny in Leuconostco cenos
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Molecule type: protein
A;Residues: 1-5 ARE>
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C;Species: Leuconostoc oenos phage P54
C;Date: 31-Mar_1993 #sequence_revision 22-May-1998 #text_change 22-May-1998
                                                   T-cell receptor beta chain V-D-J region (141-1CO) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jul-1992 #text_change 30-May-1997
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8.7%; Score 2; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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8.7%; Score 2; DB 2; Length 5;
Best Local Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                              A;Status: translation not shown
A;Molecule type: mRNA
A;Residues: 1-5 <FEE.
A;Experimental source: day 19 fetal thymus, strain BALB/c
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R;Arendt, E.K.; Lonvaud, A.; Hammes, W.P.
J. Gen. Microbiol. 137, 2135-2139, 1991
A;Title: Lysogeny in Leuconostoc oenos.
A;Reference number: A44817; MUID:92085033; PMID:1748868
A;Accession: F44817
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RESULT 69
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Gaps

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Copper resistance protein A - Pseudomonas syringae pv. tomato (fragment)
C,Species: Pseudomonas syringae pv. tomato
C,Species: 19-Unr-1992 #sequence_revision 19-Jun-1992 #text_change 24-Jun-1993
C,Accession: A41225
R;Cha, J.S.; Cooksey, D.A.
Proc. Natl. Acad. Sci. U.S.A. 88, 8915-8919, 1991
A,Title: Copper resistance in Pseudomonas syringae mediated by periplasmic and outer mem A;Accession: A41225
A;Accession: A41225
A;Accession: A41225
A;Accession: Afl225
A;Accession: Afl225
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A;Accession: Afl225
A;Accession: Afl225
A;Accession: Techniminary
A;Acsions: 1-5 < CHA>
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Best Logal Similarity 0.0%; Pred. No. 2.8e+05;
Matches 0; Conservative 0; Mismatches 1; Indels
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Search completed: February 12, 2003, 10:52:50 Job time : 17 secs

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GenCore version 5.1.3
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OM protein - protein search, using sw model

Run on:

US-09-403-440A-1 23

1 FXXLW 5 Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

112892 segs, 41476328 residues Searched:

Total number of hits satisfying chosen parameters:

20

Minimum DB seq length: 5 Maximum DB seq length: 5

Post-processing: Minimum Match 0%

Maximum Match 100% Listing first 100 summaries

SwissProt_40:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Result No.	Score	% Query Match	% Query Match Length DB	DB	OI OI		Description	ion
1	11	47.8			BPP7 B(BOTIN	P30425 bo	bothrops in
0	11	47.8	ß	ч	UF01 M	MOUSE	P38639 m	mus musculu
٣	9	26.1	ιΩ	_	AL14 C	CARMA	P81817 C	carcinus ma
4	9	26.1	S	Н	EI03 L	LITRU	P82099 1	litoria rub
ហ	9	26.1	ស	Н	EIO4 LITRU	ITRU	P82100 1	litoria rub
9	9	26.1	Z,	-	FARP ARTTE	RITR	P41853 a	artioposthi
7	9	26.1	S	-	PAP2 PARM	ARMA	P81864 p	pardachirus
80	9	26.1	Ŋ	Н	RE11 L	LLITRU	P82070 1	litoria rub
6	9	26.1	S	-	RE21_L	ITRU	P82071 1	litoria rub
10	9	26.1	S	Н	RE31 L	ITRU	P82072 1	itoria rub
11	9	26.1	Ŋ	~1	RE32 L	ITRU	P82073 1	itoria rub
12	9	26.1	Ŋ	Н	SUGA_ACHDO	СНДО	P19991 a	acheta dome
13	9	26.1	S	Н	TPISC	ANFA	P54714 C	canis famil
14	9	26.1	Ŋ	-	UC22 MAIZE	AIZE	P80628 Z	zea mays (m
15	4	17.4	'n	٦	Ŀ	PERAM	P01373 p	periplaneta
16	4	17.4	ß	٦	PSK DAUCA	UCA	P58261 d	daucus caro
17	7	8.7	2	Н	BIOA C	CITFR	P13071 C	citrobacter
18	7	8.7	2	П	BIOB CITFR	ITFR	P12997 C	citrobacter
19	0	0.0	5	П	TRM3 E	ECOLI		escherichia
20	0	0.0	ĸ	٦	UXA4 C	CHLTR	P38005 C	chlamydia t

ALIGNMENTS

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01-APR-1993 (Rel. 25, Created)
01-FBB-1994 (Rel. 28, Last sequence update)
01-FBB-1994 (Rel. 28, Last annotation update)
Bradykinin-potentiating peptide 55,2 (5A) (Angiotensin-converting
                           5 AA.
                           STANDARD;
                           BOTIN
                          BPP7 BOT
P30425;
RESULT 1
BPP7 BOTIN
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P81817; 30-MAY-2000 (Rel. 39, Created) 30-MAY-2000 (Rel. 39, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update)

STANDARD;

AL14 CARMA

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Cintra A.C.O., Vieira C.A., Giglio J.R.;

"Primary structure and biological activity of bradykinin potentiating peptides from Bothrops insularis snake venom.";

J. Protein Chem. 9:221-227(1990).

-!- FUNCTION: THIS PEPTIDE BOTH INHIBITS THE ACTIVITY OF THE ANOIOTENSIN-CONVERTING BUXYME AND ENHANCES THE ACTION OF BRADYKININ BY INHIBITING THE KINASES THAT INACTIVATE IT.

IT ACTS AS AN INDIRECT HYPOTENSIVE AGENT.

PIR; G37196; G37196.
                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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MEDLINE=95009907; PubMed=7523108;

MEDLINE=95009907; PubMed=7523108;

MEDLINE=95009907; PubMed=7523108;

MEDLINE=95009907; PubMed=7523108;

"Separation and sequencing of familiar and novel murine proteins using preparative two-dimensional gel electrophoresis.";

Blectrophoresis 15:735-745(1994).

-I MISCELLANEOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN PROTEIN IS: 6.6, ITS MW IS: 19 kDa.
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
Bothrops insularis (Island jararaca) (Queimada jararaca).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Lepidosauria; Squamata; Scleroglossa; Serpentes; Colubroidea;
Viperidae; Crotalinae; Bothrops.
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01-0CT-1994 (Rel. 30, Last sequence update)
01-0CT-1995 (Rel. 31, Last annotation update)
Unknown protein from 2D-page of fibroblasts (P19) (Fragment).
                                                                                                                                                                                                                                                                                                                    1 1 PYRROLIDONE CARBOXYLIC ACID
5 AA; 629 MW; 776DC37326B00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                   47.8%; Score 11; DB 1; Length 5; 100.0%; Pred. No. 1.1e+05; vative 0; Mismatches 0; Indels
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                                                                                                                          TISSUE=Venom;
MEDLINE=90351557; PubMed=2386615;
                                                                                                                                                                                                                                                                                                     Hypotensive agent; Venom.
MOD RES 1 1
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                                                                                                                                                                                                                                                                                                                                                                                                      1; Conservative
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P38639;
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ID A114 CP
AC P818I7;
DT 30-MAY-
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DT 30-MAY-
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Litoria rubella (Desert tree frog).
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P81864;
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Carcinus maenas (Common shore crab) (Green crab).

Bukaryota, Metazoa, Arthropoda; Mandibulata; Pancrustacea; Crustacea;

Malacostraca; Bumalacostraca; Bucarida; Decapoda; Pleccyemata;

Brachyura; Bubrachyura; Portunoidea; Portunidae; Carcinus.

NCBI_TaxID=6759;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Litoria rubella (Desert tree frog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria.
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                                                                                                                                                                                                                                                                 "Isolation and identification of multiple neuropeptides of the
                                                                                                                                                                                                                                                                                    allatostatin superfamily in the shore crab Carcinus maenas.";
Eur. J. Biochem. 250.727-734 (1997).
-1- FUNCTION: MAY ACT AS A NEUROTRANSMITTER OR NEUROMODULATOR.
-1- SIMILARITY: BELONGS TO THE ALLATOSTATIN FAMILY.
Neuropeptide; Amidation; Multigene family.
5 S AMIDATION (POTEWILL).
SEQÜENCE 5 AA; 586 MW; 672879D5AB300000 CRC64;
                                                                                                                                                           TISSUE=Cerebral ganglion, and Thoracic ganglion;
MEDLINE=98121193; PubMed=9461295;
Duve H., Johnsen A.H., Maestro J.-L., Scott A.G., Jaros P.P.,
Thorpe A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
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5 AA; 630 MW; 668761F2C9A00000 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
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15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
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-!- SUBCELLULAR LOCATION: SECRETED.
MOD_RES
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nes 1; Conserva
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P82099;
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P82100;
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Platyhelminthes; Turbellaria; Seriata; Tricladida;
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                                                                                                                                                                                                                                                                                                                                                                     TISSUE=Skin secretion;
Wabhitz P.A., Bowle J.H., Tyler M.J., Wallace J.C.;
Wabhitz P.A., Bowle J.H., Tyler M.J., Wallace J.C.;
Wabhitz From the skin glands of the Australian buzzing tree frog
Litori electrica. Comparison with the skin peptides from Litoria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             30-MAY-2000 (Rel. 39, Created)
30-MAY-2000 (Rel. 39, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
16-OCT-2001 (Rel. 40, Last annotation update)
Pardachirus marmoratus (Red sea moses sole).
Pardachirus marmoratus (Red sea moses sole).
Actinopterygii, Mebazoa, Chordata; Craniata; Vertebrata; Buteleostomi;
Actinopterygii, Neopterygii, Teleostei; Buteleostei; Soleoidei; Soleidae; Pardachirus.
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria.
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Maule A.G., Shaw C., Halton D.W., Curry W.J., Thim L.;
"RYIRFamide: a turbellarian FMRFamide-related peptide (FaRP).";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE 5 AA; 616 MW; 61F2D1A059A00000 CRC64;
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SEQUENCE 5 AA; 754 MW; 69D4004B44600000 CRC64;
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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Terricola, Geoplanidae, Arthurdendyus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Aust. J. Chem. 52:639-645(1999).
-1- SUBCELLULAR LOCATION: SECRETED.
Amphibian skin; Amidation.
**AMIDAT**
**A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRT;
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                                                                                                                                                                            NCBI_TaxID=104895;
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us-09-403-440a-1.closed.rsp

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15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                       NCBI_TaxID=104895;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Amphibian skin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Rubellidin 3.1.
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ID RE31 LITRU
AC P82072;
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RE32_LITRU
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                                                           TISSUE=SKin secretion;
MEDLINE=1057369; PubMed=3782138;
Lazarovici P., Primor N., Loew L.M.;
Luzarovici P., Primor N., Loew L.M.;
"Purification and pore-forming activity of two hydrophobic polypeptides from the secretion of the Red sea moses sole (Pardachirus marmoratus)."

Biol. Chem. 261:16704-16713(1986).

--- FUNCTION: EXHIBITS UNUSUAL SHARK REPELLENT AND SURFACTANT PROPERTIES. FORMS VOLTAGE-DEPENDENT, ION-PERMEABLE CHANNELS IN MEMBRANES. AT HIGH CONCENTRATION CAUSES CELL MEMBRANE LYSIS.

--- SUBUNIT: MONOWER. IN AQUEDOUS SOLUTION EXISTS AS A TETRAMER.

--- SUBCELLULAR LOCATION: Secreted.

--- SUBLELLULAR LOCATION: Secreted.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The structure of new peptides from the Australin red tree frog Litoria rubella. the skin peptide profile as a probe for the study of evolutionary trends of amphibians.",
Aust. J. Chem. 49:955-963 (1996).
-!- FUNCTION: SHOW NEITHER NEUROPEPTIDE ACTIVITY NOR ANTIBIOTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Gaps
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Litoria rubella (Desert tree frog).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi, Amphibia, Batrachia, Anura; Neobatrachia, Bufonoidea, Hylidae, Pelodryadinae, Litoria.
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Steinborner S.T., Wabhitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Tyler M.J., Wallace J.C.;
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-1- MASS SPECTROMETRY: MW=598; METHOD=FAB.
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                                                                                                                                                                                                                                                                                                                                                                       5 5 AA; 614 MW; 7769C9C9C8100000 CRC64;
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15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
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(Rel. 41, Last sequence update)
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NCBI TaxID=31087;
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P82071;
15-JUN-2002
15-JUN-2002
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P82070:
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SEQUENCE
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Best Local (
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DT 18-JUN
DT 18-JUN
DT 15-JUN
DE RUBATY
OC AMBLIC
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TISSUE-Skin secretion;
Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Tryler M.J., Walled J.C.,
"The structure of new peptides from the Australin red tree frog
Lintoria rubella'. the skin peptide profile as a probe for the study
of evolutionary trends of amphibians.";
Aust. J. Chem. 49:955-963(1996).
-!- FUNCTION: SHOW NEITHER NEUROPEPTIDE ACTIVITY NOR ANTIBIOTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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Aust. J. Chem. 49:955-963(1996).

-!- FUNCTION: SHOW NEITHER NEUROPEPTIDE ACTIVITY NOR ANTIBIOTIC
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                                                                          Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Amphibia; Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae;
Pelodryadinae, Litoria.
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Sukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria
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Steinborner S.T., Wabnitz P.A., Waugh R.J., Bowie J.H., Gao C.,
Tyler M.J., Wallace J.C.;
Tyler M.J., Wallace J.C.;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 26.1%; Score 6; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 1.1e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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        MOD_RES
        5
        AMIDATION.

        SEQUENCE
        5 AA;
        656 MW;
        71A9C9CB10300000 CRC64;

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15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Rubellidin 2.1.
Litoria rubella (Desert tree frog)
                                                                                                                                                                                                                                                                  SEQUENCE, AND MASS SPECTROMETRY.
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(TIM) (Fragment)

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15-JUL-1998 (Rel. 36, Last annotation update) Triosephosphate isomerase (BC 5.3.1.1) (TIM)
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SEQUENCE
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Bukaryota; Metazoa; Arthropoda; Mandibulata; Pancrustacea; Hexapoda;
Insecta; Pterygota; Neoptera; Orthopteroidea; Orthoptera; Ensifera;
Gryllidae; Gryllinae; Acheta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Gaps
                                                                                                                                                                                                                                                                                                              0; Gaps
                                                                                                                                                                Wahnitz P.A., Bowie J.H., Tyler M.J., Wallace J.C., Bowie from "Peptides from the skin glands of the Australian buzzing tree frog Litori electrica. Comparison with the skin peptides from Litoria
                           15-JUN-2002 (Rel. 41, Created)
15-JUN-2002 (Rel. 41, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2002 (Rel. 41, Last annotation update)
Rubellidin 3.2.
Litoria rubella (Desert tree frog).

Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Pelodryadinae; Litoria.

NCBI_TaxID=104895;
                                                                                                                                                                                                  rubella.";
Aust. J. Chem. 52:639-645(1999).
-!- FUNCTION: SHOW NEITHER NEUROPEPTIDE ACTIVITY NOR ANTIBIOTIC
-ACTIVITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN DORSAL GLANDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Wicker C., Wicker C.;
"Isolation and structure of a peptide isolated from the suboesophageal ganglion of Acha domesticus (orthoptera)."; Comp. Biochem. Physiol. 88c:185-187(1987).
-: SUBCELLULAR LOCATION: MAIN PEPTIDE FROM THE SUBOESOPHAGEAL
                                                                                                                                                                                                                                                                                       26.1%; Score 6; DB 1; Length 5; ilarity 100.0%; Pred. No. 1.1e+05; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                   5 AA; 570 MW; 71A9C9C862A00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                       01-FEB-1991 (Rel. 17, Created)
01-FEB-1991 (Rel. 17, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                   5 AA.
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(Rel. 34, Last sequence update)
         5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Suboesophageal ganglion pentapeptide.
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         PRT;
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         STANDARD;
                                                                                                                                                      rissuE=Skin secretion;
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                                                                                                                                                                                                                                                        Amphibian skin.
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ID TPIS'CANFA

AC P5471-1996 (

DT 01-OCT-1996 (
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P19991;
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         RE32 LITRU
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                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                             RESULT 12
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Towart P., Riccardi F., Morin C., Damerval C., Huet J.-C.,
Pernollet J.-C., Zivy M., de Vienne D.;
"The maize two dimensional gel protein database: towards an integrated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, PACC clade,
Panicoideae, Andropogoneae, Zea.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                   Theor. Appl. Genet. 93:997-1005 (1996).
-!- MISCELLANGOUS: ON THE 2D-GEL THE DETERMINED PI OF THIS UNKNOWN PROTEIN IS: 6.1, ITS MW IS: 30.4 kDa.
Maize-2DPAGE; P80628; COLEOPTILE.
                                                                                                                                                                                                                                                                                                                                                        -!- PATHWAY: PLAYS AN IMPORTANT ROLE IN SEVERAL METABOLIC PATHWAYS. -!- SUBUNIT: HOMODIMER (BY SIMILARITY). -!- SIMILARITY: BELONGS TO THE TRIOSEPHOSPHATE ISOMERASE FAMILY. HSC-2DPAGE; P54714; DOG.
                      Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia; Eutheria, Carnivora, Fissipedia, Canidae, Canis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR000652; Tripphos ismrse.
PROSITE; PS0171; TIM; PARTIAL.
Isomerase; PS0171; TIM; PARTIAL.
Pentose shunt.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .,
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15-JUL-1999 (Rel. 38, Last annotation update)
Unknown protein from 2D-page of etiolated coleoptile (Spot 474)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ·.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26.1%; Score 6; DB 1; Length 5; 100.0%; Pred. No. 1.18+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           26.1%; Score 6; DB 1; Length 5; 100.0%; Pred. No. 1.1e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 AA; 654 MW; 72CB19C9C0300000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 AA; 550 MW; 64444862C9A00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 AA.
                                                                                                                                                             TISSUE=Heart;
MEDLINE=98163340; PubMed=9504812;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-OCT-1996 (Rel. 34, Created)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STANDARD;
Canis familiaris (Dog)
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EMBRYOS.
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ID BIOA_CITFR
AC P13071;
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SEQUENCE
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                                             beta)]
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                                                                                                           Limulus polyphemus (Atlantic horseshoe crab), and Carcinus maenas (Common shore crab) (Green crab).

Eukaryota, Metazoa, Arthropoda; Mandibulata; Pancrustacea; Hexapoda; Insecta; Pterygota; Neoptera; Orthopteroidea; Dictyoptera; Blattaria; Blattoidea; Blattidae; Periplaneta.

NCBI_TaxID=6978, 6850, 6759;
                                                                                                                                                                            SPECIES=P.americana;
MEDLINE=76074708; PubMed=576;
Starratt A.N., Brown B.E.;
"Structure of the pentapeptide proctolin, a proposed neurotransmitter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                            Stangier J., Dircksen H., Keller R., "Identification of proctolin in
                                                                                                                                                                                                                                               SPECIES=P.americana;
MEDILINES B1225865; PubMed=6113690;
O'Shea M., Adams M.E.;
"Pencapeptide (proctolin) associated with an identified neuron.";
Science 213:567-569(1981).
                                                                                                                                                                                                                                                                                                                                                                                                                            Identification of proctolin in the central nervous system of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
0
                                                                                                                                                                                                                                                                                                         SPECIES=L.polyphemus;
MEDLINE=90287800; PubMed=2356151;
Groome J.R., Tillinghast E.K., Townley M.A., Vetrovs A.,
Watgon W.H. III, Hunt D.F., Griffin P.R., Alexander J.E.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Length 5; . 1.1e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 AA; 649 MW; 71B7673B44600000 CRC64;
                                                                      21-JUL-1996 (Rel. 01, Last sequence update) 01-FEB-1995 (Rel. 31, Last annotation update) Proctolin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 AA.
                                                    5 A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       17.4%; Score 4; DB 1
100.0%; Pred. No. 1.1
ive 0; Mismatches
                                                                                                    Periplaneta americana (American cockroach)
                                                                                                                                                                                                                                                                                                                                                           horseshoe crab, Limulus polyphemus.";
Peptides 11:205-211(1990).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRT;
                                                                                                                                                                                                                                                                                                                                                                                           SPECIES=C.maenas;
MEDLINE=86232789; PubMed=2872661;
                                                                    Created)
                                                                                                                                                                                                                      Life Sci. 17:1253-1256(1975)
                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                   (Rel. 01,
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Matches 1; Conserv
                                                                                                                                                                                                                                       BIOLOGICAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Neuropeptide.
                                                                   21-JUL-1986
                                                                                                                                                                                                                                                                                                                                           Shabanowitz
                                                  PRCT PERAM P01373;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 16
PSK_DAUCA
ID _ PSK_DAUCA
                                                                                                                                                                                                               insects.
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Query Match
                                  RESULT 15
PRCT_PERAM
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16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Phytosulfokine-alpha (PSK-alpha) [Contains: Phytosulfokine-beta (PSK-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PATHWAY: Bictin biosynthesis.
SUBUNIT: HOMODIMER.
SIMILARITY: BELONGS TO CLASS-III OF PYRIDOXAL-PHOSPHATE-DEPENDENT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        "Transcriptional regulation and gene arrangement of Escherichia coli, Citrobacter freundii and Salmonella typhimurium biotin operons."; Gene 67:203-211(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- SUBCELLULAR LOCATION: Secreted.
-!- PTM: SULFATION IS IMPORTANT FOR ACTIVITY AND FOR THE BINDING TO PUTATIVE MEMBRANE RECEPTOR (BY SIMILARITY).
-!- SIMILARITY: BELONGS TO THE PHYTOSULFOKINE FAMILY.
                                                                                                                                                                                                                                                                                                                                                              SEQUENCE, AND IDENTIFICATION BY MASS SPECTROMETRY.
STRAIN=cv. US-Harumakigosun;
MEDLINE=20212743; PubMed=10750705;
Hanai H., Matsuno T., Yamamoto M., Matsubayashi Y., Kobayashi T.,
Kamada H., Sakagami Y.,
"A secreted peptide growth factor, phytosulfokine, acting as a
stimilatory factor of carrot sometic embryo formation.";
Plant Cell Physiol. 41:27-32 (2000).
-!- FUNCTION: IN PRESENCE OF 2,4-D, STIMULATES PROLIFERATION OF THE
CELLS, BUT DOES NOT STIMULATE DIFFERENTIATION INTO THE SOMATIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-JAN-1990 (Rel. 13, Created)
01-JAN-1990 (Rel. 13, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Adenosylmethionine-8-amino-7-oxononanoate aminotransferase
(EC 2.6.1.62) (7,8-diamino-pelargonic acid aminotransferase) (DAPA
                                                                                                                                                                                                Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, eudicotyledons, core eudicots,
Asteridae, euasterids II, Apiales, Apiaceae, Daucus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CATALYTIC ACTIVITY: S-adenosyl-L-methionine + 8-amino-7-oxononanoate = S-adenosyl-4-methylthio-2-oxobutanoate + 7,8-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3 3 SULFATION.
5 AA; 687 MW; 76C1BB504B300000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PHYTOSULFOKINE-BETA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 1.1e+05;
2; Mismatches C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SULFATION
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MEDLINE-89006280; PubMed=2971595;
Shiuan D., Campbell A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COFACTOR: PYRIDOXAL PHOSPHATE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17.48;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      aminotransferase) (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Growth factor; Sulfation.
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nes 0; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
                                                                                                                                               beta)].
Daucus carota (Carrot).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AMINOTRANSFERASES
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                                                                                                                                                                                                                                                                                                      NCBI_TaxID=4039;
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Shiuan D., Campbell A.;
"Transcriptional regulation and gene arrangement of Escherichia coli,
"Citrobacter freundii and Salmonella typhimurium biotin operons.";
Gene 67:203-211(1988).
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                                                                                                                                                                                                                                                                                                                                                        0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -i- PATHWAY: Biotin biosynthesis; last step.
-i- SIMILARITY: BELONGS TO THE BIOTIN AND LIPOIC ACID SYNTHETASES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-JAN'1990 (Rel. 13, Created)
01-JAN-1990 (Rel. 13, Last sequence update)
15-JUN-2002 (Rel. 14, Last annotation update)
Biotin synthase (EC 2.8.1.6) (Biotin synthetase) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity 0.0%; Pred. No. 1.18+05; nes 0; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                             Score 2; DB 1; Length 5; Pred. No. 1.1e+05; 1; Mismatches 0; Indelt
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     8.7%; Score 2; DB 1; Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 - CATALYTIC ACTIVITY: Dethiobiotin + sulfur = biotin.
                                                                                                                                                      EMBL; M21922; -; NOT_ANNOTATED_CDS.
InterPro; IPR000954; Aminotran 3.
PROSITE; PS00600; AA TRANSFER CLASS 3; PARTIAL.
Biotin blosynthesis, Transferase; Aminotransferase;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NON_TER 5 5 S SEQUENCE 5 AA; 532 MW; 75A5B1EDD6F00000 CRC64;
                                                                                                                                                                                                                                                       NON TER 5 5 SEQUENCE 5 AA; 582 MW; 6AAABIBIA6F00000 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EMBL; M21922; -; NOT ANNOTATED CDS.
Biotin biosynthesis; Iron-sulfur; Transferase.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 AA.
                                                                                                                                                                                                                                                                                                                 8.7%;
0.0%; P
                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 0.00,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
                                                                                                                                                                                                                    Biotin Diosymmers.
Pyridoxal phosphate.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  BIOB CITFR
                                                                                                                                                                                                                                                                                                                                                                                              4 L 4
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RESULT 19

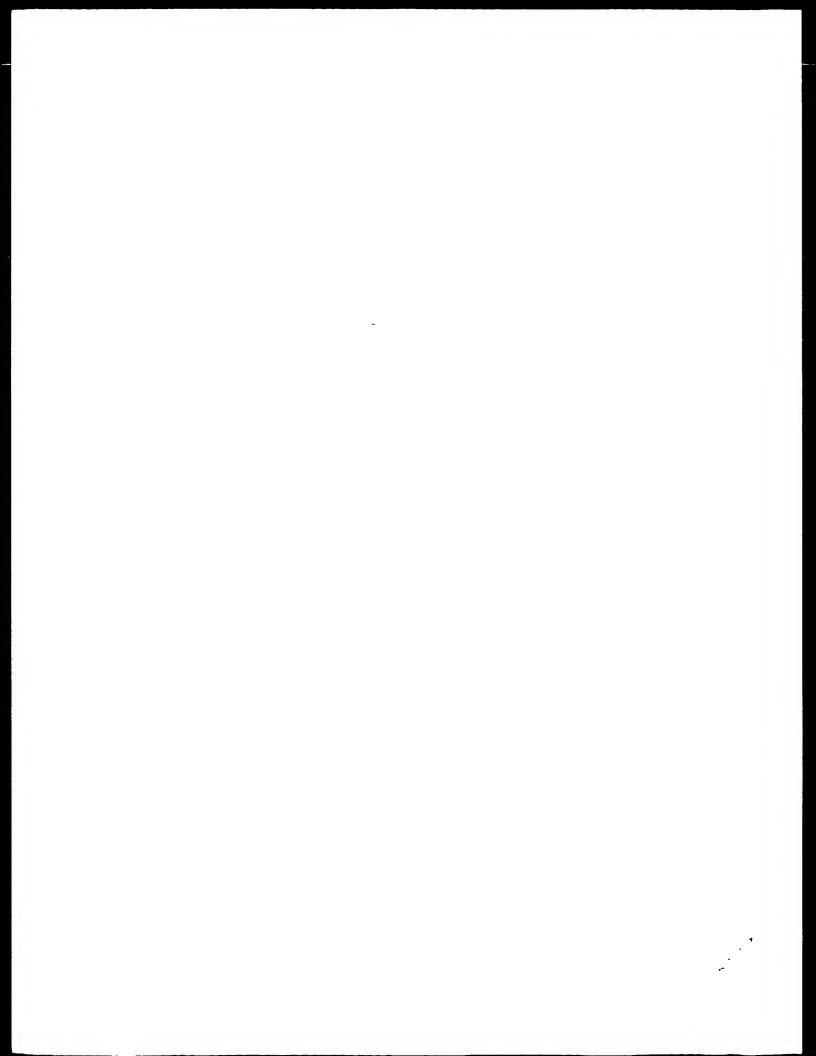
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                 MEDINE-88227859; bubMed=2836369;
MEDINE-88227859; bubMed=2836369;
Inamoro S., Yoshioka Y., Ohtsubo B.;
Indentification and characterization of the products from the trady and trary genes of plasmid R100.";
and trary genes of plasmid R100.";
J. Bacteriol. 170:2749-2757(1988).
I. FUNCTION: TRANSFER GENE PROTEIN. IS INVOLVED IN THE CONJUGATION PROCESS OF BACTERIAL GELLS FOR THE EXCHANGE OF PLASMID DNA.
I. SUBCELLULAR LOCATION: Cytoplasmic.
I. SIMILARITY: TO TRAM PROTEIN OF OTHER PLASMIDS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRAIN=L2/434/Bu;
Bini L., Santucci A., Magi B., Marzocchi B., Sanchez-Campillo M.,
Comanducci M., Christianen G., Birkelund S., Vtretou E., Ratti G.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           -i- MISCELLANEOUS: ON THE ZD-GEL THE DETERMINED PI OF THIS UNKNOWN PROTEIN IS: 4.5, ITS MW IS: 28 kDa.
Siena-ZDPAGE; P38005; -.
NON TER
                                                                                                                                            Plasmid IncFII R100.
Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
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01-OCT-1994 (Rel. 30, Last sequence update)
01-OCM-1994 (Rel. 39, Last annotation update)
Unknown protein from 2D-page from elementary body (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.0%; Score 0; DB 1; Length 5; Best Local Similarity 0.0%; Pred. No. 1.1e+05; Matches 0; Conservative 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 AA; 474 MW; 75BAA865AA800000 CRC64;
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Bacteria; Chlamydiales; Chlamydiaceae; Chlamydia.
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                                 01-JAN-1990 (Rel. 13, Created)
01-JAN-1990 (Rel. 13, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; M20941; -; NOT_ANNOTATED_CDS.
PIR; A32014; A32014.
Conjugation; Plasmid; DNA-binding.
NON_TER
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STANDARD;
                                                                                         TraM protein (Fragment).
                                                                                                                              Escherichia coli.
                                                                                                                                                                                                    NCBI_TaxID=562;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI TaxID=813;
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1D UXA4 CHLTR
                                                                                                                                                                                  Escherichia
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 TRM3 ECOLI
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Best Loćal Similarity 0.0%; Pred. No. 1.1e+05; Matches 0; Conservative 0; Mismatches 1; Indels 0; Gaps 0;

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Search completed: February 12, 2003, 10:51:53 Job time : 12 secs



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Gaps .,

0; Indels

Searched:

Sequence:

Run on:

5 AA.

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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Archosauria, Aves, Neognathae, Galliformes, Phasianidae, Phasianinae,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hordeum vulgare (Barley).
Eukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Magnoliophyta, Liliopsida, Poales, Poaceae, Pooideae,
Triticeae, Hordeum.
NCBI_TaxID=4513;
                                                                                                                        PubMed=613771;
Dockray G.J., Reeve J.R. Jr., Shively J., Gayton R.J., Barnard C.S.;
M.A novel active pentapepide from chicken brain identified by
antibodies to FMRPamide.";
                                                                                                                                                                                     Nature 305:328-330(1983).
-!- FUNCTION: MAY FUNCTION AS A NEUROTRANSMITTER OR MODULATOR.
-!- SIMILARITY: BELONGS TO THE FARP (FMRFAMIDE RELATED PEPTIDE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bacteria; Firmicutes; Bacillus/Clostridium group; Bacillales; Bacillacea; Bacillus.
NCBL_TaxID=1396;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 8.7%; Score 2; DB 2; Length 5; Best Local Similarity 0.0%; Pred. No. 6.7e+05; Matches 0; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                   h 26.1%; Score 6; DB 13; Length 5; Similarity 100.0%; Pred. No. 6.7e+05; 1; Conservative 0; Mismatches 0; Indels
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Last sequence update)
Last annotation update)
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01-OCT-2001 (TrEMBLrel. 18, Last sequence update)
01-OCT-2001 (TrEMBLrel. 18, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 AA; 623 MW; 6B01AAA336F00000 CRC64;
                                                                                                                                                                                                                                                                     S AA; 645 MW; 69D4073767400000 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             88 kDa protein (Fragment).
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Gallus gallus (Chicken).
                                                                                          SEQUENCE, AND SYNTHESIS
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                                                                                                                                                                                                                                                                                                                Local Similarity
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                                                         NCBI_TaxID=9031;
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                                                                                                                                                                                                                                                    Neuropeptide.
SEQUENCE 5
                                                                                                           TISSUE=BRAIN;
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P83073 bacillus ce
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                                                                                                        February 12, 2003, 10:48:54 ; Search time 28 Seconds (without alignments) 36.794 Million cell updates/sec
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              GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd.
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P83308;
01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
FMRFamide-like neuropeptide (LFLRF-amide).
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                                                                            OM protein - protein search, using sw model
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P83073
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2: sp_bacteria:*
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Gapop 10.0 , Gapext 0.5
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RESULT 1 P83308

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Hydrolase; Glycosidase; Carbohydrate metabolism; Seed; Germination;
Calcium; Multigene family.
NON TER
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SEQUENCE 5 AA; 600 MW; 61E3344DD6F00000 CRC64;
STRAIN=HIMALAYA; TISSUE=ALEURONE LAVER;

MEDLINE=91329704; PubMed=1831055;

"Gacobsen J.V., Clase T.J.;

"Control of transient expression of chimaeric genes by gibberellic acid and abscisic acid in protoplasts prepared from mature barley alucrone layers.",

"Analyzic Acily177: ENDOHYDROLYSIS OF 1,4-ALPHA-GLUCOSIDIC LINKAGES IN OLIGOSACCHARIDES AND POLYASACCHARIDES.

"-- COPACTOR: BINDS A CALCIUM ION REQUIRED FOR ITS ACTIVITY.

BARLEY.

BARLEY.
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8.7%; Score 2; DB 10; Length 5;
Best Local Similarity 0.0%; Pred. No. 6.7e+05;
Matches 0; Conservative 1; Mismatches 0; Indels
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Perfect score:

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Sequence:

Scoring table:

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Endothelin metal c
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TAN-1462A - cyclic
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EGIII-like cellula
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p53/MDM2 binding i
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| SIDSZ/gcgdata/geneseqfyemeseqgyembl/AA2002.DAT:*
| SIDSZ/gcgdata/geneseqfyemeseqgyembl/AA2002.DAT:*
| SIDSZ/gcgdata/geneseqfyemeseqgyembl/AA2002.DAT:*
                                                                                                                  February 12, 2003, 10:45:58 ; Search time 80 Seconds (without alignments) 8.328 Million cell updates/sec
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             GenCore version 5.1.3
(c) 1993 - 2003 Compugen Ltd.
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AAB07280
AAW17808
AAK89911
AAK89912
AAK58280
AAC20822
AAR29438
AAR29448
                                                                                                                                                                                                                                                                      BLOSUM62
Gapop 10.0 , Gapext 0.5
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Match Length
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Maximum |
Listing
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82.6
82.6
82.6
82.6
82.6
73.9
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Score

Result No.

Endothelin recepto	HBV surface antige	Core polypeptide f	Peptide #36 for il	Viral DP178/107-li													
AAY97920	AAY97921	AAY97922	AAY97935	AAY97936	AAY97937	AAY97950	AAY97951	AAY97952	AAY85099	AAY85100	AAY85101	AAY85102	AAY85103	AAY89380	AAM47406	ABB00788	
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9	9	9	9	9	9	9	9	Ģ	9	ő	9	9	9	9	9	9	
12	15	15	15	15	15	15	15	15	15	15	15	15	15	15	15	15	
84	85	98	87	88	83	90	91	92	93	94	9	96	97	86	66	100	

ALIGNMENTS

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This invention describes a novel method for preparing a peptide or recombinant antibody, which mimics an antigenic carbohydrate. The peptides and recombinant antibodies prepared to mimic antigenic carbohydrates can be used to enhance binding of anti-antigenic carbohydrates antibodies to the antigenic carbohydrate, e.g. as vaccine carbohydrate antibodies to the antigenic carbohydrate, e.g. as vaccine adjuvants. The peptides can be used to inhibit binding of a ligand to a receptor, which is an antigenic carbohydrate. The methods are used to prepare the peptides and antigenic carbohydrates. Carbohydrate-conjugate vaccines could be used in tumour immunity and cancer therapy. The peptides and antibodies can also be used as antibacterials. Peptides that mimic carbohydrate antigens can be carbohydrates and partigens can be used to develop a longer lasting immune response. Other advantages of the peptide mimics are; (1) the chemical composition and purity of synthesized peptides can be precisely defined; (2) the immunogenicity of the peptides can be significantly enhanced by polymerization or addition or relatively small carrier molecules that reduce the total
                                                                                                                                                                                        Mimotope, antigen, carbohydrate, antibody, mimic, vaccine, adjuvant, tumour immunity, cancer therapy, antibacterial, immune response, immunogenicity, anti-idiotype, T cell response manipulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptides and recombinant antibody mimics of carbohydrate antigens,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                used for, e.g. treatment of cancer and infection
                                                                                                                                               Carbohydrate antigen peptide mimotope 11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 21; Page 72; 88pp; English.
                  AAY33038 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                          99WO-US02405.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   98US-0073690.
                                                                                                      (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              (UYPE-) UNIV PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1999-527317/44.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Kieber-Emmons T;
                                                                                                                                                                                                                                                                                                                        WO9940433-A1.
                                                                                                                                                                                                                                                                                                                                                                                                          04-FEB-1999;
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                                                                                                      03-NOV-1999
                                                                                                                                                                                                                                                                                                                                                                 12-AUG-1999.
                                                             AAY33038;
AAY33038
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DB 21; Length 5;

87.0%; Score 20;

Sequence Query Match

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Tumour metastasis requires detachment of malignant cells from the primary tumour, penetration of blood or lymph vessels and attachment to the endothelium of distant organs, ultimately resulting in the formation of new tumours. The selectin family of adhesion molecules is implicated in this process. E-selectin family of adhesion molecule six implicated in this process. E-selectin is a calcium-dependent molecule expressed by activated vascular endothelium. E-selectins bind to glyoconjugates carrying a terminal tetrasaccharide Lewis antigen, which are found on tumour cell surfaces. One such Lewis antigen is Lewis Y (LeY). The binding of selectin molecules to their ligands is thought to be an important step in metastasis. Therefore, inhibition of E-selectin-dependent carbohydrate—mediated interactions is thought to be a target for anti-cancer therapy. The present sequence is a non planar-X-planar type motif. This motif is thought to bind to anti-LeY antibody BRS5-2. BRS5-2 is a peptido-mimetics of E-selectins. Peptides that block E-selectin-LeY binding and therefore adhesion of tumour cells and leukocytes to endothelial cells inhibit metastasis.
                                                                                                                                                                                                                                       0
amount of antigen required for immunization; (3) peptide synthesis may be more practical than synthesis of carbohydrate-protein conjugates or the production of anti-idiotypes; (4) peptide mimicking sequences can be engineered into DNA plasmids for DNA vaccination to further manipulate T cell responses. ANY33028-Y333087 represent the carbohydrate antigen peptide mimotopes described in the invention.
                                                                                                                                                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptidomimetics of carbohydrate Lewis ligands useful for modulating
                                                                                                                                                                                                                                       0;
                                                                                                                                                                                              87.0%; Score 20; DB 20; Length 5; 60.0%; Pred. No. 7.8e+05; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Human; peptido-mimetic; tumour metastasis; E-selectin; adhesion molecule; Lewis antigen; anti-adhesion therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Motif binding to anti-Lewis antigen anibody BR55-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inflammation, metastasis and angiogenesis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (WIST-) WISTAR INST ANATOMY & BIOLOGY. (UYPE-) UNIV PENNSYLVANIA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Example 5; Page 37; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                           AAB07280 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US26277.
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                                                                                                                                                                                                                                       Conservative
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                                                                                                                                                                                                           Local Similarity
les 3; Conserv
                                                                                                                                                           5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Unidentified.
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                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                              ovulation; menstrual cycle; oestrus; meat quality; growth; spawning;
fish; testosterone; superagonist; leuprorelin acetate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           'note= "N-pToluenesulphonylarginine, D-form residue"
                                                                                                                                                                                                                                                                                                                                Luteinising hormone releasing hormone receptor; LH-RH; antagonist; sex-hormone; human; veterinary medicine; cancer; prostate; uterus; breast; pituitary; prostatomegaly; endometriosis; hysteromyoma;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cyclic penta:peptide(s), some new, as LH-RH receptor antagonists - used to treat or prevent sex hormone dependent disorders, e.g. cancer, also for control of pregnancy and menstruation and to
                                                                                                                                                                                                                                                                                                                                                                                         precocious puberty; amenorrhoea; premenstrual syndrome; cyclic; multilocular ovarian syndrome; comedo; pregnancy; contraception; ovulation; menstrual cycle; oestrus; meat quality; growth; spawn.
                        ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'note= "Joined via a peptide linkage to Trp5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "Joined via a peptide linkage to Phel"
                                                                                                                                                                                                                                                                                             Cyclic pentapeptide #24 used in LH-RH receptor antagonist.
Pred. No. 7.8e+05;
0: Mismatches 2; Indels
                       0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 44; Page 172; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                  AAW17808 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Furuya S, Kato K, Kitada C;
 60.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    95JP-0110933.
95JP-0106775.
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                                                                                                                                                                                                                                                             (first entry)
                     3; Conservative
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Best Local Similarity
Matches 3; Conserv
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Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New peptides of the invention which interfere with the binding of the p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein binding motif between amino acids 16-33 (AAR89914). The peptides were identified by modifying the p53 consensus binding sequence by substitution of an Alanine at each pos. and identifying which amino acid changes altered binding to expressed MDM2. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                        Score 19; DB 17; Length 5;
Pred. No. 7.8e+05;
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100.0%; Pred. No. 7.8e+05;
iive 0; Mismatches 0;
                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                   /note= Any amino acid
                                                                                                                                                                                                                                                                                                                                                                           'note= Any amino acid
                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                             AAR89911 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Claim 4; Page 29; 46pp; English.
                                                                                                                                                                                                                                                        p53/MDM2 binding inhibitor #1.
                                    82.68;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     95US-0424957.
94US-0277660.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             95WO-GB01719,
                     Query Match
Query Match
Best Local Similarity 60.00,
                                                                                                                                                                                                                              10-SEP-1996 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Lane DP, Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1996-105905/11.
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Best Local Similarity
             5 AA;
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Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New peptides of the invention which interfere with the binding of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 82.6%; Score 19; DB 17; Length 5;
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                                                                                                                                                                                                                                            /note= Any amino acid
                                                                                                                                                                                                               Location/Qualifiers
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                         AAR89912 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 5; Page 29; 46pp; English.
                                                                                                             p53/MDM2 binding inhibitor #2.
                                                                                                                                                                                                                                                                                                                                95WO-GB01719.
                                                                                                                                                                                                                                                                                                                                                           95US-0424957.
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                                                                                  10-SEP-1996 (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                    (UYDU-) UNIV DUNDEE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1996-105905/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    other malignancies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                         20-JUL-1994;
                                                                                                                                                                                                                                                                                                                                20-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                           19-APR-1995;
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                                                                                                                                                                                                                                                                                                   01-FEB-1996
                                                                                                                                                                                   Synthetic.
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                                                      AAR89912;
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RESULT 5
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Human p53 consensus hydrophobic interaction motif.

27-MAR-2000 (first entry)

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receptor function, for example NRS can stimulate transcription in response to hormone binding by recruiting coactivator proteins to the propose to hormone binding by recruiting coactivator proteins to the cativity of a transcription domain. AF2, that is part of the cativity of a transcription domain. The method may be used to identify an agonist or antagonist of coactivator binding to a nuclear receptor. The methods can also be used to identify an agonist or antagonist of coactivator binding to a nuclear receptor of interest. The methods are applicable to generating new compounds that distinguish nuclear receptor isoforms. This can facilitate generation of either tissue specific or function-specific compounds. Nuclear receptors have been implicated in a variety of medical disorders, including breast cancer, prostate cancer, cardiac arrhythmia, infertility, osteoporosis, Ohyperthyroidism, hypercholesterolaemia and obbsity. Sequences AMYSB282. AMYSB302 represent the regions (NR-boxes 1-3) of pl60 coactivator binding proteins which interact with the nuclear receptor coactivator binding
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The invention relates a method of identification of a compound that modulates coactivator binding to a nuclear receptor. The method compounds that fit spatially into a nuclear receptor (NR) coactivator binding site of interest using an atomic structural model of a nuclear receptor coactivator binding site. The ceptor compounds are screened by their ability to bind to an NR coactivator binding site, leading to the identification of a compound which modulates coactivator binding. Nuclear receptors are a superfamily of hormone/ ligand activated transcription factors, and includes thyroid hormone receptors (TRS), retinoid receptors (RARs and RARS), peroxisome receptors (PRARs and XPARS), vientin D receptors (VDRS), oestrogen receptors (ERS), glucocorticoid receptors (GRS), progestin receptors (RRS), mineralocorticoid receptors (MRS), and and solved receptors (RRS) and solved r
Nuclear receptor; pl60 coactivator; NR-box; coactivator binding domain; NR; AF2 transactivation domain; wild-type; thyroid hormone receptor; TR; retinoid receptor; RAR; RXR; peroxisome receptor; PPAR; RXR; peroxisome receptor; PR; yitamin D receptor; VDR; oestrogen receptor; ER; glucocorticoid receptor; GR; progestin receptor; PR; mineralocorticoid receptor; MR; androgen receptor; AR; Grip-1; Tif2; NcoA-2; RAC3; AIB1; TRAM-1; P/CIP; SRC1; breast cancer; prostate cancer; cardiac arrhythmia; infertility; osteoporosis; hyperthyroidism; hypercholesterolaemia; obesity;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Identifying modulators of nuclear receptor coactivator binding useful for generating new compounds which distinguish nuclear receptor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fletterick R, Kushner PJ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'note= "Xaa = any amino acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "Xaa = any amino acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Example 14B; Page 12; 281pp; English.
                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Darimont B, Feng W, F
West BL, Yamamoto KR;
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                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     30-MAR-1999;
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                                                                                                                                                                                                                                                                                                                                    Homo sapiens
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Wagner RL,
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sites (AAYS8307-Y58328). The NR coactivator binding sites comprise two regions of the NR; helices 3-6 (H3-H6), and helix 12 (H12). The coactivator NR-boxes all share a consensus sequence LXXLL (AAYS8278), with the NR-box 2 motifs sharing the consensus motif ILXXLL (AAYS8279). Sequences AAYS8280-Y58281 represent the hydrophobic interaction motifs of p53 and VP16 used in an exemplification of the invention. Sequences AAYS8329-Y58240 represent mutant versions of the human Grip-1 NR-box 2 sequence (AAYS8283), and sequences AAYS8341-Y58353 represent mutant versions of the human Grip-1 NR-box 2 versions of NR coactivator binding sites, used in exemplifications of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cytostatic; cardiovascular; antiarrhythmic; vasotropic; osteopathic; endocrinal; antilipaemic; anorectic; coactivator binding; lead compound; nuclear receptor; NS; peptidomimetic; scaffold; breast cancer; nuclear receptor-based disorder; prostate cancer; cardiac arrhythmia; infertility; osteoporosis; hyperthyroidism; hypercholesterolaemia; obesity; ExxLW motif; p53; glucocorticoid; GR; protein co-ordinate data.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Kushner PJ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Method for identifying agonist and antagonist compounds used for modulating nuclear receptor coactivator binding, which are useful treating nuclear receptor-based disorders e.g. breast cancer, infertility, or hyperthyroidism -
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, Geistlinger TR, Arnol
                                                                                                                                                                                                        82.6%; Score 19; DB 21; Length 5; 100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                            100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hydrophobic interaction FxxLW motif of p53.
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/note= "Xaa is unknown"
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Yamamoto KR,
                                                                                                                                                                                                                                                                                                                                                                         AAO20822 standard; Peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             29-JUN-2001; 2001WO-US20969.
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,, West BL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 2002-315164/35.
                                                                                                                                                                                                                 Local Similarity
                                                                                                                                                                         5 AA;
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                                                                                                                                                                                                                                                                 1 FXXLW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                        08-JUL-2002
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                                                                                                                                            invention.
                                                                                                                                                                           Sequence
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                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Guy RK,
                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                             RESULT 7
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The invention relates to a method for identifying a compound that

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modulates coactivator binding to a nuclear receptor (NR). The method comprises modelling test compounds that fit electrostatically into an NR caactivator binding site (BS) using an appropriate atomic structural model of (BS); screening the test compounds in an assay characterised by binding of a test compound to (BS); and identifying a library or test compound that modulates coactivator binding to the NR. The methods of the invention are useful for identifying peptides, peptidomimetics, or small netral or synthetic organic molecules that modulate nuclear receptor natural or synthetic organic molecules that modulate nuclear receptor activity, particularly new lead compounds, scaffolds and combinatorial libraries. The methods are used in selecting and characterising structure function relationships of natural or synthetic coactivator compounds, which are useful in treating nuclear receptor-based disorders e.g. breast cancer, prostate cancer, cardiac arrhythmia, infertility, osteoporosis, hyperchycidism, hypercholesterolaemia or obesity. This sequence compended (RR) nuclear receptor of the invention is able to interact with peptides containing this motif.
                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hypertension, myocardial infarction; congestive heart failure; endotoxic shock; subarachnoid haemorrhage; asthma; arrhythmias; acute renal failure; preeclampsia; diabetes; metabolic; endocrinological; neurological; disorders.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The peptide is an endothelin antagonist useful in controlling hypertension, myocardial infarction, congestive heart failure,
                                                                                                                                                                                                                                                                                                                                                                               Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                          82.6%; Score 19; DB 23; I
100.0%; Pred. No. 7.8e+05;
ive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR29438 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endothelin antagonist peptide.
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                                                                                                                                                                                                                                                                                                                                                                                                                  5; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                           5 AA;
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               acute renal failure, preeclampsia, diabetes and metabolic, endocrinological and neurological disorders. Administration is oral parenteral or by inhalation in doses of 0.01-20 (6sp. 0.01-10) mg/kg/day. It may be prepared by conventional peptide synthesis.
                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                 Hypertension; myocardial infarction; congestive heart failure; endotoxic shock; subarachnoid haemorrhage; asthma; arrhythmias; acute renal failure; preeclampsia; diabetes; metabolic; endocrinological; neurological; disorders.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New peptide(s) used as endothelin antagonists - for treating hypertension, metabolic and endocrine disorders, heart failure, diabetes, asthma, neurological disorders, etc.
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endotoxic shock, subarachnoid haemorrhage, asthma, arrhythmias,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The peptide is an endothelin antagonist useful in controlling
                                                                                                                             Query Match 73.9%; Score 17; DB 13; Length 5; Best Local Similarity 40.0%; Pred. No. 7.8e+05; Matches 2; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Taylor MD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers
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                                                                                                                                                                                                                                                                                                           AAR29448 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                         Endothelin antagonist peptide.
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                                                                                                  5 AA;
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1 FDIIW 5
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endocrine disorders, congestive heart failure, myocardial infarction, endotoxic shock, subarachnoid haemorrhage, arrhythmia, asthma, acute and chronic renal failure, preeclampsia, diabetes, neurological disorders, pulmonary hypertension, ischaemic disease, ischaemic bowel disease, gastric mucosal damage, Raynaud's disease, restenosis, percutaneous transluminal coronary angioplasty, angina and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Novel antagonists of endothelin are claimed which are C-terminal hexapeptides and analogs of BT-1. The first (N-terminal) amino acid of the new peptides has D-configuration. The peptides are claimed generically. The present peptide is a specifically claimed example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New hexa:peptide derivs. inhibiting endothelin - for treatment of e.g. renal failure, hypertension, asthma, restenosis, angina, cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      of the generic compounds.
The peptides are useful for treating hypertension, metabolic and
                                                                                                                                                                                          Endothelin C-terminal peptide analog, useful as antagonist.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           He JX, Taylor MD;
                                                                                                                                                                                                                        Endothelin; ET-1; receptor; antagonist.
                                                                                                                                                                                                                                                                                 Location/Qualifiers
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                                                                                                    AAR69220 standard; peptide; 5 AA.
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es 2; Conserv
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                                                                                                                                                                                                                                                                                                Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                     17-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 FLDIW 5
1 FXXLW 5
                           S
                                                                                                                                                                                                                                                                                                                                                                         07-JUL-1994.
                                                                                                                                                               06-MAR-1995
                         1 FLDIW
                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR69260;
                                                                                                                                  AAR69220;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 11
                                                                    RESULT 10
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0; Gaps

Score 17; DB 13; Length 5; Pred. No. 7.8e+05; 1; Mismatches 2; Indels

73.9%;

Query Match
Best Local Similarity 40.0°
Matches 2; Conservative

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(WARN ) WARNER LAMBERT
                                                                                                                                      WPI; 1994-234617/28
                                                                                                                                                                                                                                                                                                                                                                       Local Similarity
les 2; Conserv
                                                                                                                                                                                                                                                                                                                                            5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified-site
                                                            17-DEC-1993;
                                                                                                                                                                                                                                                                                                                                                                                                   1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                     1 FDIIW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       08-AUG-1990;
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                        WO9414843-A.
                                          07-JUL-1994.
                                                                                                                                                                   e.g. renal f
cancer etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-OCT-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-JUN-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic.
                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR13339;
                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cyclic.
                                                                                                                                                                                                                                                                                                                                                                         Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR13339
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                                                                                                                                                                                                                                                                                                                                                        The peptides are useful for treating hypertension, metabolic and endocrine disorders, congestive heart failure, myocardial infarction, endocraic shock, subarachnoid haemorrhage, arrhythmia, asthma, acute and chronic renal failure, precalampsia, diabetes, neurological disorders, pulmonary hypertension, ischaemic disease, ischaemic bowel disease, gastric mucosal damage, Raynaud's disease, restenosis, percutaneous transluminal coronary angioplasty, angina and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                           Novel antagonists of endothelin are claimed which are C-terminal hexapeptides and analogs of BT-1. The first (N-terminal) amino acid of the new peptides has D-configuration. The peptides are claimed generically. The present peptide is a specifically claimed example
                                                                                                                                                                                                                                                   New hexa:peptide derivs. inhibiting endothelin - for treatment of e.g. renal failure, hypertension, asthma, restenosis, angina,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  _note= "Ac-D-Phe or Ada-D-Phe, where Ada is
                  Endothelin C-terminal peptide analog, useful as antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Endothelin C-terminal peptide analog, useful as antagonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 17; DB 15; Length 5;
Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2; Indels
                                                                                                                                                                                                               Taylor MD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                                                                                                                               не лх,
                                      Endothelin; ET-1; receptor; antagonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ET-1; receptor; antagonist
                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                                                                                                                                                                                                                                                        Claim 5; Page 120; 146pp; English.
                                                                                               /note= "Ac-D-Phe"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR69210 standard; peptide; 5 AA.
                                                                                                                                                                                                              Depue P, Doherty AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                         73.9%;
                                                                                                                                                       93WO-US12377
                                                                                                                                                                        92US-0995480
07-MAR-1995 (first entry)
                                                                                                                                                                                          (WARN ) WARNER LAMBERT CO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     06-MAR-1995 (first entry
                                                                                                                                                                                                                                                                                                                                                  of the generic compounds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                 WPI; 1994-234617/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Best_Local Similarity
Matches 2; Conserv
                                                                                    Misc-difference 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Key
Misc-difference 1
                                                                                                                                                     17-DEC-1993;
                                                                                                                                                                       21-DEC-1992;
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1 FDFIW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 FXXLW 5
                                                                                                                                  07-JUL-1994
                                                                                                                WO9414843-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Endothelin;
                                                                                                                                                                                                                                                                       cancer etc.
                                                         Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                             Cody WL,
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR69210;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR69210
 113X8XX8X1X1
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generically. The present peptide is a specifically claimed example of the generic compounds.

The peptides are useful for treating hypertension, metabolic and endocrine disorders, congestive heart failure, mycoardial infarction, endotoxic shock, subarachnoid haemorrhage, arrhythmia, asthma, acute and chronic renal failure, preeclampsia, diabetes, neurological disorders, pulmonary hypertension, ischaemic disease, ischaemic bowel disease, gastric mucosal damage, Raynaud's disease, restenosis, percutaneous transluminal coronary angioplasty, angina and cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Novel antagonists of endothelin are claimed which are C-terminal hexapetides and analogs of ET-1. The first (N-terminal) amino acid of the new peptides has D-configuration. The peptides are claimed generically. The present peptide is a specifically claimed example
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New hexa:peptide derivs. inhibiting endothelin - for treatment of e.g. renal failure, hypertension, asthma, restenosis, angina,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            73.9%; Score 17; DB 15; Length 5; llarity 40.0%; Pred. No. 7.8e+05; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Taylor MD;
1-adamantyl-acetyl"
                                                                                                                                                                                                                                                                                                                                                                                                                                                   не лх,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Endothelin antagonist peptide WS 7338.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /label= D-aile, D-val
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 5; Page 118; 146pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR13339 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                               Cody WL, Depue P, Doherty AM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'label= D-Trp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /label= D-Glu
                                                                                                                                                                                                                   93WO-US12377.
                                                                                                                                                                                                                                                                                              92US-0995480
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Gaps

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Length 5;
                                                                        0; Indels
isolated from cultures of Streptomyces sp. no. 7338.
See also AAR13339 and AAR13909.
                                                   65.2%; Score 15; DB 12; L
100.0%; Pred. No. 7.8e+05;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                         Location/Qualifiers
                                                                                                                                                                  AAR13909 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; page 1; 15pp; Japanese.
                                                                                                                                                                                                                             Endothelin antagonist peptide
                                                                                                                                                                                                                                                                                                            'label= D-Val
                                                                                                                                                                                                                                                                                                                                 /label= D-Trp
                                                                                                                                                                                                                                                                                                                                                    /label= D-Glu
                                                                                                                                                                                                                                                                                                                                                                                                                                  89GB-0019726.
                                                                                                                                                                                                                                                                                                                                                                                                                 90JP-0212952.
                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                      (FUJI ) FUJISAWA PHARM KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                          2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        endothelin antagonist
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 1991-241129/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
                                                              Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 AA;
                                                                                                                                                                                                                                                                                                  Modified-site
                                                                                                                                                                                                                                                                                                                       Modified-site
                                                                                                                                                                                                                                                                                                                                          Modified-site
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                                                                                                                                                                                                          22-OCT-1991
                                                                                                                                                                                                                                                                      Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 LW 5
                                                                                              4 LW 5
                                                                                                                2 LW 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 LW 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                      AAR13909;
                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
                                                    Query Match
                                                                                                                                                                                                                                                  Cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 16
AAR38437
                                                                          Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                              RESULT 15
                                                                                                                                                         AAR13909
                                                                                                                                                                            Вb
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     SSXS
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0
                                                                                                                                                                                                                                         0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The peptide is prepd. by cyclisation of the synthetic peptide. The Trp residue at posn. 3 may be side chain protected with a lower alkanoyl gp; the Glu residue at posn. 4 may be protected with an al (lower) alkyl gp. An analogous natural peptide, WS 7338, may be
                                                                                                                                  The Trp residue at posn. 3 may be side chain protected with a lower alkanoyl gp; the Glu residue at posn. 4 may be protected with an al (lower) alkyl gp. An analogous natural peptide, WS 7338, may be isolated from cultures of Streptomyces sp. no. 7338.
                                                                                                                             peptide is prepd. by cyclisation of the synthetic peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New peptide prepd. by culturing Streptomyces spp - used as endothelin antagonist.
                                                                          New peptide prepd. by culturing Streptomyces spp - used as endothelin antagonist.
                                                                                                                                                                                                                  65.2%; Score 15; DB 12; Length 5; ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                   AAR14025 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 1; page 1; 15pp; Japanese.
                                                                                                        Claim 1; page 1; 15pp; Japanese.
                                                                                                                                                                            See also AAR13909 and AAR14025.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            'label= D-aIle
                                                                                                                                                                                                                                                                                                                                                                                              Endothelin antagonist peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /label= D-Trp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /label= D-Glu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               90JP-0212952.
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                89GB-0019726.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (FUJI ) FUJISAWA PHARM KK.
                                                                                                                                                                                                                                                                                                                                                                           (first entry)
                                   (FUJI ) FUJISAWA PHARM KK.
                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1991-241129/33.
                                                       WPI; 1991-241129/33.
                                                                                                                                                                                                  5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-AUG-1990;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   31-AUG-1989;
                31-AUG-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            04-JUN-1991.
                                                                                                                                                                                                                                                                                                                                                                           22-OCT-1991
                                                                                                                                                                                                                                                              4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                2 LW 3
                                                                                                                                                                                                   Sequence
                                                                                                                                                                                                                                                                                                                                                      AAR14025;
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                                                                                                                                              The peptide is prepd. by cyclisation of the synthetic peptide. The Trp residue at posn. 3 may be side chain protected with a lower alkanoyl gp; the Glu residue at posn. 4 may be protected with an al (lower) alkyl gp. An analogous natural peptide, WS 7338, may be isolated from cultures of Streptomyces sp. no. 7338.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .
0
peptide prepd. by culturing Streptomyces spp - used as
                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                     65.2%; Score 15; DB 12; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
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WPI; 1993-282913/36.
   Misc-difference 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference 2
                                                                                                                                                                                                                                                                                                             5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference 3
                     Misc-difference
                                                Misc-difference
                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference
                                                                                                JP05194589-A
                                                                                                                                     30-JUN-1992;
                                                                                                                                                       31-JUL-1991;
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                                                                                                                  03-AUG-1993
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                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                    4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR38436;
                                                                                                                                                                                                                                                                                                                                                                                       4 LW
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                     The sequences given in AAR38436-38 are cyclic peptides which act as endothelin receptor antagonists. They may be used in the treatment of cardiac disease, acute renal insufficiency or asthma.
                                                                                                                                                                                                                                                                                                                                             Novel cyclic peptide(s) used as endothelin receptor antagonists - for treating cardiac diseases, e.g. myocardial infarction, acute renal insufficiency or asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .
0
                                                                                                                                                             /label= OTHER
/note= "alpha-Abu, delta-Ava or epsilon-Aca"
                                                                         Cyclic, endothelin receptor, antagonist, cardiac disease, myocardial infarction, acute renal insufficiency, asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             65.2%; Score 15; DB 14; Length 5; ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cyclic, endothelin receptor; antagonist; cardiac disease; myocardial infarction; acute renal insufficiency; asthma.
                                                      Endothelin receptor antagonist peptide #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Endothelin receptor antagonist peptide #3.
                                                                                                                                            /note= "D-form residue"
                                                                                                                                                                                          'note= "D-form residue"
                                                                                                                                                                                                             /note= "D-form residue"
                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                    Example 4-6; Page 6; 8pp; Japanese.
AAR38437 standard; peptide; 5 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR38438 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                     92JP-0173065
                                                                                                                                                                                                                                                                                        91JP-0191454
                                                                                                                                                                                                                                                                                                         (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                    07-MAR-1994 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity
                                                                                                                                                                                                                                                                                                                            WPI; 1993-282913/36.
                                                                                                                                                     Misc-difference 2
                                                                                                                                                                               Misc-difference 3
                                                                                                                                Misc-difference
                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                             5 AA;
                                                                                                                                                                                                                                                                    30-JUN-1992;
                                                                                                                                                                                                                                                                                       31-JUL-1991;
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                                                                                                                                                                                                                                                  03-AUG-1993
                                                                                                      Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   4 LW 5
                  AAR38437;
                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 LW S
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Gaps
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                                                                                                                                                                                                                                                                                                                                                  Novel cyclic peptide(s) used as endothelin receptor antagonists for treating cardiac diseases, e.g. myocardial infarction, acute renal insufficiency or asthma
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/note= "alpha-Abu, delta-Ava or epsilon-Aca"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65.2%; Score 15; DB 14; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cyclic, endothelin receptor; antagonist, cardiac disease, myocardial infarction; acute renal insufficiency; asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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'note= "D-form residue"
                                                                                                                 /note= "DOform residue"
                                                                                'note= "D-form residue"
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                                                                                                                                                                                                                                                                                                                                                                                                                       Disclosure; Page 6; 8pp; Japanese.
                               /label= OTHER
/note= "beta-Asp"
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Best Local Similarity 100.0
Matches 2; Conservative
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action of endothelin. They are useful as drugs for improving circulatory function in mammals, in treatment of myocardial infarction, acute renal dysfunction or asthma. Thesse peptides may be prepared by conventional methods of solid or aqueous phase
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                                                                                                  Best Local Similarity 100.0 Matches 2; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference 3
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                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference 5
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                                                                    5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    asthma, etc.
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                                               synthesis.
                                                                                                                                                                                                                                                                                                                                       Synthetic.
                                                                                                                                    4 LW 5
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                                                                                          Query Match
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                                                                                                                                                          The sequences given in AAR38436-38 are cyclic peptides which act as endothelin receptor antagonists. They may be used in the treatment of cardiac disease, acute renal insufficiency or asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                        Cyclic, endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
                                                                                           Novel cyclic peptide(s) used as endothelin receptor antagonists -
for treating cardiac diseases, e.g. myocardial infarction, acute
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                                                                                                                                                                                                                            Length 5;
                                                                                                                                                                                                                           65.2%; Score 15; DB 14; Length 5; 100.0%; Pred. No. 7.8e+05; trive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                    Endothelin receptor inhibitor peptide #1.
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                                                                                                                                       Example 1-3; Page 6; 8pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                   AAR35418 standard; peptide; 5 AA.
                                                                                                                   renal insufficiency or asthma
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                           91JP-0191454.
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     92JP-0173065.
                                                (TAKE ) TAKEDA CHEM IND LTD
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Best Local Similarity 100.0
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
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                                                                      WPI; 1993-282913/36
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                                                                                                                                                                                                      5 AA;
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     30-JUN-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cyclic; endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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65.2%; Score 15; DB 14; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 2; Conservative 0; Mismatches 0;
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Matches
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                                                                                                                       Cyclic; endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
                                                                                                                                                                                                                                                                                                                                                                                       The sequences given in AAR35418-28 are cyclic peptides which act as endothelin inhibitors. They strongly inhibit the vasoconstrictive action of endothelin. They are useful as drugs for improving circulatory function in mammals, in treatment of myocardial infarction, acute renal dysfunction or asthma. Thesse peptides may be prepared by conventional methods of solid or aqueous phase
                                                                                                                                                                                                                                                                                                                                 New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           65.2%; Score 15; DB 14; Length 5; 100.0%; Pred. No. 7.8e+05; Live 0; Mismatches 0; Indels
                                                                                                       Endothelin receptor inhibitor peptide #4.
                                                                                                                                                                                 /note= "D-form residue"
                                                                                                                                                                                                                   /note= "D-form residue"
                                                                                                                                                                                                  'note= "D-form residue'
                                                                                                                                                                  Location/Qualifiers
                                                      AAR35421 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                          Example 4; Page 8; 8pp; Japanese.
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                                                                                       03-AUG-1993 (first entry)
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Matches 2; Conserv
                                                                                                                                                                         Misc-difference 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 AA;
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                                                                                                                                                 Synthetic.
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                                                                        AAR35421;
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               3 LW
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                                        RESULT 21
AAR35421
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Gaps
                                                     Cyclic; endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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100.0%; Pred. No. 7.8e+05;
iive 0; Mismatches 0;
Endothelin receptor inhibitor peptide #5.
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                                                                                                                                                                                                                                     'note= "D-form residue"
                                                                                                                                                                                                                                                                                     'note= "D-form residue"
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                                                                                                                                                                                  Location/Qualifiers
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                                                                                                                                                                                                        Misc-difference 1
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                                                                                                                                Synthetic
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RESULT 25
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Cyclic; endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
                                                                                                                                  New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
                                                                                                                                                                                                                                                                                               .<u>;</u>
                                                                                                                                                                                                                                                                              Length 5;
                                                                                                                                                                                                                                                                             65.2%; Score 15; DB 14; Length 5; llarity 100.0%; Pred. No. 7.88+05; Conservative 0; Mismatches 0; Indels
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                 /note= "D-form residue"
/note= "D-form residue"
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                                                                                    91JP-0220610.
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                                                                                                    (TAKE ) TAKEDA CHEM IND LTD
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                                                                    30-AUG-1991;
                                                                                    30-AUG-1991;
                                   JP05059098-A
                                                   09-MAR-1993
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                                                                                                                                                                                                                                                                                                            The sequences given in AAR35418-28 are cyclic peptides which act as endothelin inhibitors. They strongly inhibit the vasoconstrictive action of endothelin. They are useful as drugs for improving circulatory function in mammals, in treatment of myocardial infarction, acute renal dysfunction or asthma. Thesse peptides may be prepared by conventional methods of solid or aqueous phase synthesis.
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                                                                                                        New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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(TAKE ) TAKEDA CHEM IND LTD.
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endothelin inhibitors. They strongly inhibit the vasoconstrictive action of endothelin. They are useful as drugs for improving circulatory function in mammals, in treatment of myocardial infarction, acute renal dysfunction or asthma. Thesse peptides may be prepared by conventional methods of solid or aqueous phase synthesis.
                                                                                                                                                                                                                                                                                              Cyclic, endothelin, inhibitor, vasoconstriction, drug, circulation, mammal, myocardial infarction; renal dysfunction, asthma.
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                                                                                           65.2%; Score 15; DB 14; Length 5; 100.0%; Pred. No. 7.8e+05; Ative 0; Mismatches 0; Indels
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                                                                                                                 Conservative
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                                                                                                       Local Similarity
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                                                                                                                                                                                                                                                                                                                                               Cyclic, endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
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100.0%; Pred. No. 7.8e+05;
iive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Endothelin; metal conjugate; complex; diagnostic; contrast agent;
X-ray imaging; NMR imaging; pathological vascular change; stenolic;
atherosclerosis; radiopharmaceutical; non-invasive diagnosis;
arterial wall lesion; pharmacokinetic.
                                      Cyclic; endothelin; inhibitor; vasoconstriction; drug; circulation; mammal; myocardial infarction; renal dysfunction; asthma.
                                                                                                                                                                                                                                                                                   New cyclic penta:peptide derivs. contg. one L-aminoacid - inhibit endothelin receptor activity, vasoconstriction etc., useful for treating myocardial infarction, acute renal dysfunction and
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                  Endothelin receptor inhibitor peptide #11.
                                                                                                        /note= "D-form residue"
                                                                                                                                             /note= "D-form residue"
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                                                                                     Location/Qualifiers
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Matches 2; Conserv
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This invention describes novel complexes of compounds which have the formula B-L-(K)b with metal ions of atomic number 21-32, 37-39, 42-51 and 57-83 where E = residue of an endothelin (or its derivative, partial sequence, analogue or antagonist), including forms with free SH groups that bind directly to metal, optionally labelled with radioactive iodine isotopes, L = bond, Z1-R-Z2 or a group (i), R = 1-20. Optionally constituted by 10 and/or epoxy, and coptionally interrupted by 1 or more 0, S, CO, NCO or NH (the last 2 optionally interrupted by 1 or more 0, S, CO, NCO or NH (the last 2 optionally nubstituted by 1-6C alkyl), Z1 and Z = 0, S, COO, NH, CS, NH, NCO, CSO, NH, NHO or NHCS, s and t = 0-3, B = phenyl or cyclohexyl ring, b = 0 or 1, K is a chetating residue. The compounds of the invention are useful as diagnostic agents, including contrast agents for X-ray and NMR imaging of pathological vascular changes e.g.

CX ray and NMR imaging of pathological vascular changes before these become standiopharmaceuticals. The invention allows the marking of possible early non-invasive diagnostic agents they have better pharmacokinetic properties and provide improved contrast. They have better pharmacokinetic properties and provide improved contrast. They have better pharmacokinetic properties and provide improved contrast. They have excellent in vitro and in vivo stability of sequence is a specific claimed example of a beptide used in the construction of the metalcomplex-conjugates of the invention.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            New complexes of derivatised endothelin or related cpds. - with radioactive or paramagnetic metal ions, useful as diagnostic agents for imaging pathological changes in blood vessels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Kramp W;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Hilger CS,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (DINK/) DINKELBORG L.
(UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                                                   'note= "Residue is alloDIle"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gries H, Hilger Ct
Speck U, Reiser JH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity 100.0%; Fred. No. 7.8 tes 2; Conservative 0; Mismatches
                                                                                                                                                                           /note= "D-form residue"
                                  /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DIAG-) INST DIAGNOSTIKFORSCHUNG GMBH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR47438 standard; peptide; 5 AA.
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                                                                                                                                                                                                                                                                                                                                                                                             93DE-4301871.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  93DE-4301871.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Erber S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Schier H,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           WPI; 1994-226321/28.
Misc-difference 1
                                                                                                                                         Misc-difference 5
                                                                Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Dinkelborg L,
                                                                                                                                                                                                                                                                                                                                                                                             13-JAN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-JAN-1993;
                                                                                                                                                                                                                                                    DE4301871-A1
                                                                                                                                                                                                                                                                                                                         14-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Platzek J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR47438;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
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immunosuppression; CD4; gp120; T cell activation;
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                                                                                                                                                     Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    5 AA;
                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                           10-AUG-1993;
                                                   Homo sapiens
                                                                                                                                                                                                                                                                                                          .0-AUG-1992;
                                                                                                                                                                                                                                                                                                                        12-FEB-1993;
                                                                                                                                                                                                                  WO9403487-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The peptide inhibits binding of endothelin-1 (ET) to ETA receptors and/or to ETB receptors. It can be used for detecting, distinguishing and isolating ET receptor subtypes. The peptides also act as ET antagonists and can be used to treat endothelinmediated disorders such as hypertension, pulmonary hypertension, cardiovascular disease, bronchoconstriction, asthma, inflammatory diseases, ophthalmolgic shock, anaphylactic shock, haemorrhagic menstrual disorders, obstretic conditions, erythropoietin-mediated vasoconstriction and wounds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                New cyclic peptide(s) which modulate endothelin activity - used for treating endothelin-mediated disorders such as cardiovascular
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ..
                        Cyclic; peptide; endothelin; inhibitor; receptor; detection; isolation; antagonist; hypertension; pulmonary hypertension; cardiovascular disease; bronchoconstriction; asthma; inflammatory disease; bothorhamlogic shock; anaphylactic shock; haemorrhagic shock; gastroenteric disease; renal failure; endotoxin shock; menstrual disorders; obstratic conditions; erythropoietin-mediated vasoconstriction; wound treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Monomeric peptide #1, autoimmunogenic in HIV-infected subjects.
                                                                                                                                                                                                                                           /note= "D-Trp bonded to D-Asp at position 1."
                                                                                                                                                                                'note= "D-Asp bonded to D-Trp at position
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            HIV-1; HIV-2; human immunodeficiency virus; AIDS; acquired immune deficiency syndrome; autoimmune response;
Cyclic peptide which modulates endothelin activity.
                                                                                                                                                    Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR46691 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Claim 2; Page 51; 58pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                           (IMMU-) IMMUNOPHARMACEUTICS INC.
                                                                                                                                                                                                              'label= D-His.
                                                                                                                                                                                                                                                                                                                                                                             92US-0900711
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                and respiratory diseases
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                                                                                                                                                                                              Misc-difference 3
                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                            Misc-difference
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                                                                                                                                                                                                                                                                                                                                 17-JUN-1993;
                                                                                                                                                                                                                                                                                                                                                               18-JUN-1992;
                                                                                                                                                                                                                                                                       W09325580-A
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Balaji VN,
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           New peptide(s) inducing an auto immune response in HIV infected subjects - also the multimer, proteins contg. them and derived antibodies, for treating or preventing HIV infection and for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                   /note= "C-terminus is opt. extended
by 1-5 amino acids"
                                                                                                                                                                                                                                 /note= "N-terminus is opt. extended
by 1-5 amino acids"
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                                                                                       Human Immunodeficiency Virus type 1
                                                                                                                                                                 Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           93WO-FR00803
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93FR-0001628
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therapy; prophylaxis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 1994-065602/08.
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(ZAGU/) ZAGURY J F.
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The sequences of ET-1, ET-2 and ET-3 are given in AAR60076-78. Cyclic pentapeptides, useful as endothelin receptor antagonists are given
                                                                                                                                                                                                                                                                                    0; Gaps
                                                                                                                                                         The sequences of ET-1, ET-2 and ET-3 are given in AAR60076-78. Cycl pentapeptides, useful as endothelin receptor antegonists are given in AAR60079-88. They are useful in the treatment or prophylaxis of hypertension, myocardial infarction, acute renal insufficiency or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endothelin, ET, ET-1; ET-2; ET-3; pentapeptide; receptor;
antagonist; myocardial infarction; hypertension;
acute renal insufficiency; asthma; cyclic.
                                                                                                                                                                                                                                                          / Match 65.2%; Score 15; DB 15; Length 5; Local Similarity 100.0%; Pred. No. 7.80+05; nes 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cyclic penta:peptide(s) - useful as endothelin receptor antagonists for treatment of myocardial infarction, etc.
                                                                                               Cyclic penta:peptide(s) - useful as endothelin receptor antagonists for treatment of myocardial infarction, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         rAN-1477D - cyclic ET receptor antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'note= "D-form residue"
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                                                                                                                                    Disclosure; Page 7; 13pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                  AAR60082 standard; Protein; 5 AA.
91JP-0148807.
                       91JP-0148807.
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                                               (TAKE ) TAKEDA CHEM IND LID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-MAR-1995 (first entry)
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                                                                        WPI; 1994-260512/32
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                                                                                                                                                                                                                                      5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20-JUN-1991;
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20-JUN-1991;
                       20-JUN-1991;
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                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                            Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                    Cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                The sequences of ET-1, ET-2 and ET-3 are given in AAR60076-78. Cycl pentapeptides, useful as endothelin receptor antagonists are given in AAR60079-88. They are useful in the treatment or prophylaxis of hypertension, myocardial infarction, acute renal insufficiency or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                65.2%; Score 15; DB 15; Length 5; illarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Endothelin; ET: ET-1; ET-2; ET-3; pentapeptide; receptor;
antagonist; myocardial infarction; hypertension;
acute renal insufficiency; asthma; cyclic.
                                                                                                                                                                                                                                                                                       Cyclic penta:peptide(s) - useful as endothelin receptor antagonists for treatment of myocardial infarction, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TAN-1462B - cyclic ET receptor antagonist.
                                                                        /label= alle
/note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "D-form residue"
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                                                  'note= "D-form residue"
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                          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                            Disclosure; Page 7; 13pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR60080 standard; Protein; 5 AA.
                                                                                                                                                                                      91JP-0148807.
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                                                                                                                                                                                                                                       (TAKE ) TAKEDA CHEM IND LTD.
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Rest Local Similarity ?
                                                                                                                                                                                                                                                                WPI; 1994-260512/32.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                            5 AA;
                                      Misc-difference
                                                                                                   Misc-difference
                                                              Modified-site
                                                                                                                                                                                                               20-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP06192293-A.
                                                                                                                                      JP06192293-A
                                                                                                                                                                                      20-JUN-1991;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-MAR-1995
                                                                                                                                                                12-JUL-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                              Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4 LW 5
  Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAR60080;
                                                                                                                                                                                                                                                                                                                                                                                                      asthma.
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AAR62195 standard; Protein; 5 AA.
                                                                                                                                                                                                                                                                          WPI; 1994-302689/37.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                 5 AA;
                                                               03-MAY-1995
                                                                                                                                               Homo sapiens
                                                                                                                                                                                                    10-MAR-1994;
                                                                                                                                                                WO9420141-A.
                                                                                                                                                                                  15-SEP-1994
                                                                                                                                                                                                                                                        Douvas A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Ŋ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                              AAR62195;
                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAR71907;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 LW
            36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 37
          RESULT 36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR71907
                             셤
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The sequences of ET-1, ET-2 and ET-3 are given in AAR60076-78. Cyclic
                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             pentapeptides, useful as endothelin receptor antagonists are given in AAR60079-88. They are useful in the treatment or prophylaxis of hypertension, myocardial infarction, acute renal insufficiency or
in AAR60079-88. They are useful in the treatment or prophylaxis of hypertension, myocardial infarction, acute renal insufficiency or
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0
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                                                                                                                                                                                                                           Endothelin, ET; ET-1; ET-2; ET-3; pentapeptide; receptor;
antagonist; myocardial infarction; hypertension;
acute renal insufficiency; asthma; cyclic.
                                                     Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 5;
                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cyclic penta:peptide(s) - useful as endothelin receptor antagonists for treatment of myocardial infarction, etc
                                                   65.2%; Score 15; DB 15; L
100.0%; Pred. No. 7.8e+05;
ive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           65.2%; Score 15; DB 15; L
100.0%; Pred. No. 7.8e+05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                              /note= "D-5-methyltryptophan"
                                                                                                                                                                                                         TAN-1477K - cyclic ET receptor antagonist.
                                                                                                                                                                                                                                                                                                                             'note= "D-form residue"
                                                                                                                                                                                                                                                                                                  'note= "D-form residue"
                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Disclosure; Page 7; 13pp; Japanese.
                                                                                                                                                     AAR60088 standard; Protein; 5 AA.
                                                                                                                                                                                                                                                                                                                    /label= alle
                                                                                                                                                                                                                                                                                                                                                                                                  91JP-0148807
                                                                                                                                                                                                                                                                                                                                                                                                                     91JP-0148807
                                                                                                                                                                                                                                                                                                                                                                                                                                     (TAKE ) TAKEDA CHEM IND LID
                                                                                                                                                                                         (first entry)
                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1994-260512/32.
                                                            Best Local Similarity
Matches 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                    S AA;
                                                                                                                                                                                                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 AA;
                                                                                                                                                                                                                                                                                                         Modified-site
                                                                                                                                                                                                                                                                                                                                     Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                  20-JUN-1991;
                                                                                                                                                                                         21-MAR-1995
                                                                                                                                                                                                                                                                                                                                                                                                                    20-JUN-1991;
                                                                                                                                                                                                                                                                                                                                                               JP06192293-A
                                                                                                                                                                                                                          Endothelin;
                                                                                                                                                                                                                                                                                                                                                                                 12-JUL-1994.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LW 5
                                                                                       4 LW 5
                                                                                                                                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence
                                   Sequence
                                                                                                                                                                       AAR60088;
                                                    Query Match
                                                                                                        4 LW
                  asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          asthma
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4
                                                                                                                                   RESULT 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
                                                                                                                                            AAR60088
SKGGG
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CENP-B protein; centromere protein; epitope; autoantibody; immunoinfective cluster virus; nuclear protein antigen; systemic rheumatic disorder; SRV-1 viral protease; systemic lupus erythematosus; scleroderma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A comparison of the CENP-B centromere protein sequence with proteins from immunoinfective cluster viruses revealed widespread homologies. The importance of these homologies motifs is that they are epitopes for autoantibodies occurring in high titres in systemic rheumatic disorders. Sera from such patients could be used for treatment of immunoinfective cluster virus infections, e.g. HIV-1, immunoinfective adenoviruses, human lymphotropic retroviruses,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                       CENP-B protein residues 191-195, homologous to SRV-1 protease motif.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Methods for treating immunoinfective cluster virus infections -
utilise antibodies or fragments characteristic of auto antibodies
produced by patients with rheumatic disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Endothelin cyclic peptide conjugated to chalcogen-contg. ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                rubella virus, CMV and EBV infections.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Takehana Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Page 71; 106pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR71907 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYSC-) UNIV SOUTHERN CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                    94WO-US02631.
(first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19-MAY-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Ehresmann G,
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New cyclic pentapeptides are disclosed which have a gamma-turn and a beta-turn and which are of formula Cyclo(-Al-A2-A3-A4-A5-) where the amino acids in positions 1-2-3 form a gamma-turn and the amino acids in positions 3-4-5-1 form a beta-turn in combination with the gamma-turn. Bither D-amino acid residues are chosen for Al, A3 and A5 and L-amino acids for A2 and A4; or L-amino acid residues are chosen for A1, A3 and A5 and D-residues for A2 and A5. The peptides are chosen for antagonists against the NKZ receptor or endothalin receptor. They can be used as antisathmatic, antiinflammatory or antiarthritic can be used as assay or affinity purification reagents for the
                                                                                                                                                                                                                                                                                                                                    New cyclic penta:peptide(s) having a gamma-turn and a beta-turn - used as NK2 receptor antagonists for treating asthma,
                                                                                /note= "D-Trp; and this amino acid is condensed
onto the N-terminal to form a cyclic
peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            The present sequence is a specifically claimed example of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Beta-turn; gamma-turn; NK2; neurokinin; endothelin; receptor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05;
with the C-terminal amino acid"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cyclic pentapeptide having beta-turn and gamma-turn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                antiasthmatīc; antiinflammatory; antiarthritic; affinity purification; cyclic.
                                                                                                                                                                                                                                                                                Wakimasu M;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ney Location/Qualifiers Misc-difference 1
                          /note= "Ser(Bzl)"
                                                                                                                                                                                                                                                                                                                                                                                              Claim 10; Page 16; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR69267 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                               Kikuchi T,
                                                       /note= "D-Leu"
                                                                                                                                                                                                                          93JP-0004125.
                                                                                                                                                                                                                                                     (TAKE ) TAKEDA CHEM IND LID.
                                                                                                                                                                                                94EP-0100276.
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                                                                                                                                                                                                                                                                                                                                                                  inflammation or arthritis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cyclic pentapeptides.
                                                                                                                                                                                                                                                                                                           WPI; 1994-226821/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
les 2; Conserv
                                                                                                                                                                                                                                                                                 Endo S, Inooka H,
                                        Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 AA;
                                                                    Misc-difference
              Modified-site
                                                                                                                                                                                                11-JAN-1994;
                                                                                                                                                                                                                          13-JAN-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-MAY-1995
                                                                                                                                                                     20-JUL-1994
                                                                                                                                          EP606881-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR69267;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 LW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 39
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             This peptide is one of 14 endothelin derivatives which are preferred for use in conjugates with novel bis-(N (mercapto-ethyl-or -propyl) aminocarbonylmethyl) oxide, sulphide or selenide compounds. The chalcogen-containing compounds are ligands which are complexed with a technetium or rhenium radioisotope. The iligand-peptide conjugates are useful for non-invasive, in vivo imaging, e.g. of receptors.
                                                                                                                                                                                                                                                                                                                                                                              Bis-(N-(mercapto-ethyl- or -propyl) amino:carbonyl:methyl) oxide, sulphide or selenide and derivs. - are ligands for radioactive technetium or rhenium for diagnosis and therapy, e.g. for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "D-Glu; and this amino acid is part of a
   cyclic structure formed by condensation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Beta-turn; gamma-turn; NK2; neurokinin; endothelin; receptor; antiasthmatic; antiinflammatory; antiarthritic; affinity purification; cyclic.
                    1...5
| Jabel= cyclic
/note= "attached to chalcogen-contg. ligand,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; rive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cyclic pentapeptide having beta-turn and gamma-turn.
                                                                                                                                                                                                                                                                                                                                                                                                                          non-invasive in-vivo visualisation of receptors
                                                                                                                                                                                                                                                                                               (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                                   'note= "D-form residue"
                                                                                                                           'note= "D-form residue"
                                                                                                                                                       /note= "D-form residue"
                                                          see comments"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                          Hilger C, Kramp W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR69266 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 9; Page 14; 15pp; German.
                                                                                                              label= alle
                                                                                                                                                                                                                                                                    93DE-4311022.
                                                                                                                                                                                                                                         93DE-4311022.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 1
                                                                                                                                                                                                                                                                                                                                                      WPI; 1994-311304/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 AA;
                                                                     Misc-difference
                                                                                                                                          Misc-difference
                                                                                                  Modified-site
                                                                                                                                                                                                                                                                                                                          Dinkelborg L,
                                                                                                                                                                                                                                         31-MAR-1993;
                                                                                                                                                                                                                                                                      31-MAR-1993;
                                                                                                                                                                                    DE4311022-A
                                                                                                                                                                                                             06-OCT-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR69266;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
              Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 38
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Location/Qualifiers
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                                                                                                                                                                                                                                     Endo S, Inooka H, Kikuchi T,
                                                                'note= "D-Phe"
                                                                                                                                                                                             93JP-0004125
                                                                                                                                                                                                                (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22-MAY-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                          WPI; 1994-226821/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
                                                     Misc-difference 3
                                                                          Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 AA;
    Key
Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         pentapeptides.
                                                                                                                                                                        11-JAN-1994;
                                                                                                                                                                                             13-JAN-1993;
                                                                                                                                                  20-JUL-1994.
                                                                                                                              EP606881-A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                      receptors,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR69270;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 LW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
                                                                                                                                                                                                                                                                                                                                      New cyclic pentapeptides are disclosed which have a gamma-turn and a beta-turn and which are of formula Cyclo(-A1-A2-A3-A4-A5-) where the amino acids in positions 1-2-3 form a gamma-turn and the amino acids in positions 3-4-5-1 form a beta-turn in combination with the gamma-turn. Bither D-amino acid residues are chosen for A1, A3 and A5 and L-amino acids for A2 and A4; or L-amino acid residues are chosen for A1, A3 and A5 and antagonists against the NK2 receptor or endothelin receptor. They can be used as antiasthmatic, antiinflammatory or antiarthritic agents or as assay or affinity purification reagents for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                           New cyclic penta:peptide(s) having a gamma-turn and a beta-turn used as NK2 receptor antagonists for treating asthma, inflammation or arthritis
/note= "D-Glu; and this amino acid is part of a
cyclic structure formed by condensation
with the C-terminal amino acid"
                                                                                  /note= "D-Trp; and this amino acid is condensed onto the N-terminal to form a cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Beta-turn; gamma-turn; NK2; neurokinin; endothelin; receptor;
antiasthmatic; antiinflammatory; antiarthritic;
affinity purification; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence is a specifically claimed example of the cyclic pentapeptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cyclic pentapeptide having beta-turn and gamma-turn.
                                                                                                                                                                                                                                   Wakimasu M;
                                         "Thr(Bzl)"
                                                                                                                                                                                                                                                                                                                     Claim 11; Page 16; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAR69268 standard; peptide; 5 AA
                                                              'note= "D-Leu"
                                                                                                                                                                                                                                   Inooka H, Kikuchi T,
                                                                                                                                                                                                             (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                     94EP-0100276
                                                                                                                                                                                         93JP-0004125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         22-MAY-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2; Conservative
                                           note=
                                                                                                                                                                                                                                                       WPI; 1994-226821/28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity
                                                  Misc-difference 3
                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 AA;
                               Modified-site
                                                                                                                                                                    11-JAN-1994;
                                                                                                                                                                                        13-JAN-1993;
                                                                                                                                                 20-JUL-1994
                                                                                                                            EP606881-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                   receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR69268;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                  Ś
                                                                                                                                                                                                                                   Endo
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           40
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 X S X E Z E X E X E X S X E X S X E
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beta-furn and which are of formula Cyclo(-A1-A2-A3-A4-A5-) where the amino acids in positions 1-2-3 form a gamma-turn and the amino acids in positions 3-4-5-1 form a beta-turn in combination with the gamma-turn. Bither D-amino acid residues are chosen for A1, A3 and A5 and L-amino acids for A2 and A4; or L-amino acid residues are chosen for A1, A3 and A5 and D-residues for A2 and A5. The peptides are antagonists against the NK2 receptor or endothelin receptor. They can be used as antiasthmatic, antiinflammatory or antiatthritic agents or as assay or affinity purification reagents for the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New cyclic pentapeptides are disclosed which have a gamma-turn and a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New cyclic penta:peptide(s) having a gamma-turn and a beta-turn
used as NK2 receptor antagonists for treating asthma,
inflammation or arthritis
/note= "D-Glu; and this amino acid is part of a
cyclic structure formed by condensation
with the C-terminal amino acid"
                                                                                                                                                                                                0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Beta-turn; gamma-turn; NK2; neurokinin; endothelin; receptor;
antiasthmatic; antiinflammatory; antiarthritic;
affinity purification; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 15; DB 15; Length 5; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      The present sequence is a specific example of the cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cyclic pentapeptide having beta-turn and gamma-turn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
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100.0%; Pred. No. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                New cyclic penta:peptide(s) having a gamma-turn and a beta-turn used as NK2 receptor antagonists for treating asthma,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Endothelin cyclic peptide conjugated to chalcogen-contg. ligand.
                  /note= "D-Asp; and this amino acid is part of a
cyclic structure formed by condensation
with the C-terminal amino acid"
                                                                              /note= "D-Trp; and this amino acid is condensed
  onto the N-terminal to form a cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               technetium, rhenium, in vivo imaging, atherosclerotic plaque,
bifunctional chalcogen-interrupted chelate former, cyclic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.2%; Score 15; DB 15; Length 5; ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                        The present sequence is a specific example of the cyclic
                                                                                                                                                                                                                          Endo S, Inooka H, Kikuchi T, Wakimasu M;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                         Example 5; Page 15; 29pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAR65742 standard; peptide; 5 AA.
                                                                                                  peptide"
                                                          note= "D-Leu"
                                                                                                                                                                                   93JP-0004125.
                                                                                                                                                                                                     (TAKE ) TAKEDA CHEM IND LTD.
                                                                                                                                                               94EP-0100276.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09-JUN-1995 (first entry)
                                                                                                                                                                                                                                                                                     inflammation or arthritis
                                                                                                                                                                                                                                               WPI; 1994-226821/28.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
Matches 2; Conserv
 Key
Misc-difference 1
                                                Misc-difference 3
                                                                      Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                    pentapeptides.
                                                                                                                                                               11-JAN-1994;
                                                                                                                                                                                   13-JAN-1993;
                                                                                                                        EP606881-A.
                                                                                                                                                                                                                                                                                                                                                                                                                              receptors.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR65742;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                 5
/note= "the amino group of D-Glu forms a peptide
linkage with the carboxy group of D-Trp"
                                                                                                                                                                                                                                                                                                                                                            /note= "the amino group of D-Glu forms a peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Endothelin cyclic peptide conjugated to chalcogen-contg. ligand.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bifunctional chalcogen-atom interrupted chelate formers - for radioactive isotopes, their prodn. and use in diagnostics and therapy, e.g. for non-invamive in-vivo representation of receptors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chalcogen; ligand; complex; endothelin; non-invasive imaging; technetium; rhenium; in vivo imaging; atherosclerotic plaque; bifunctional chalcogen-interrupted chelate former; cyclic.
                                                       label= cyclic
note= "attached to chalcogen-contg. ligand,
                                                                                                                                                                                                                                                                                                                                                                                         linkage with the carboxy group of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dinkelborg L, Hilger C, Kramp W, Schier H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 100.0%; Pred. No. 7.8 es 2; Conservative 0; Mismatches
                                                                                                                                                                                                                     /label= alle
/note= "D-form residue"
                                                                                                                                                                                                                                                                                                          'note= "D-form residue"
                                                                                                                                                                 /note= "D-form residue"
                                                                                                               see comments"
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR65728 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 9; Page 11; 13pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             93DE-4311023.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93DE-4311023,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-JUN-1995 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 atherosclerotic plaques.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1994-311305/39.
                                                                                                                                                                                                                                                                                Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 AA;
                                                                                                                                         Misc-difference
                                                                                                                                                                                                 Modified-site
                                                                                                                                                                                                                                                                                                                                        Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                    Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             31-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 31-MAR-1993;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        06-OCT-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DE4311023-A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 I.W
                                Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 43
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94WO-DE00369
                                                                                                                                   93DE-4311021
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                    Local Similarity 100.
                                                                                                                                                                     Dinkelborg L, Erber S,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note=
                                                                                                                                                                                      WPI; 1994-332837/41
Misc-difference 1
                                                                                                                                                                                                                                                                                                                                            5 AA;
                 Misc-difference
                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Key
Modified-site
                                                     Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               25-AUG-1995
                                                                                                                  29-MAR-1994;
                                                                                                                                  31-MAR-1993;
                                                                               WO9422491-A
                                                                                                13-0CT-1994
                                                                                                                                                                                                                                                                                                                                                                                               4 LW 5
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                                                                                                                                                                                                                                                                                                                                            Sequence
                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                                                                                                                                                                                                                                                                                                                                                               4 LW
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                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 45
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                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                        preferred for use in conjugates with novel, bifuctional chalcogen-
interrupted chelate forming compounds. The chalcogen-containing
compounds are ligands which are complexed with a technetium or
rhentium radioisotope. The ligand-peptide conjugates are useful
for non-invasive, in vivo imaging, e.g. of receptors or of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Peptide; ligand; radioactive; peptide complex; imaging; receptor; steroid; hormone; atherosclerotic plaque; targetting; endothelin; analogue; derivative; antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                          .
0
                           1..5
/label= cyclic
/note= "attached to chalcogen-contg. ligand,
see comments"
                                                                                                                                                                                                                                                                                                                This peptide is one of 14 endothelin derivatives which are
                                                                                                                                                                                                                                                          Bifunctional, chalcogen-interrupted chelate former for radioisotopes - used in diagnostics and therapy, e.g. for non-invasive in-vivo representation of receptors
                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; Ative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "Forms peptide linkage with DTrp5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cyclic peptide ligand #2 for radiopharmaceutical agent.
                                                                                                                                                                                                      (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                    note= "D-form residue
                                                                                               'note= "D-form residue
                                                                                                                /note= "D-form residue
                location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                         Kramp W,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR63183 standard; peptide; 5 AA
                                                                                                                                                                                                                                                                                             Claim 9; Page 12; 13pp; German.
                                                                                     'label= alle
                                                                                                                                                                    93DE-4310999
                                                                                                                                                                                      93DE-4310999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         16-JUN-1995 (first entry)
                                                                                                                                                                                                                         Dinkelborg L, Hilger C,
                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                     atherosclerotic plaques.
                                                                                                                                                                                                                                          WPI; 1994-311297/39.
                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                            Misc-difference
                                                                                                        Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                      5 AA
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Modified-site
                                                                                                                                                                    31-MAR-1993;
                                                                                                                                                                                     31-MAR-1993;
                                                                                                                                  DE4310999-A
                                                                                                                                                   06-OCT-1994
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Synthetic
                                                                                                                                                                                                                                                                                                                                                                                       Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                          4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                         Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                           4
                                                                                                                                                                                                                                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      44
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The sequences given in AAR73170-83 represent peptide ligands which are used with the radioactive peptide complex of the invention. The complexes may be used to image receptors, specifically of steroid hormones, receptor contg. issue and/or atherosclerotic plaque. These peptides allow the complex to be targetted to a specific tissue. These peptides are pref. derived from endothelin, and represent analogues, derivatives or antagonists.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5
/note= "optionally has a carboxyl, amido, hydrophobic or macromolecular carrier gp. attached"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 "optionally has an amino, acetyl, 9-fluorenylmethoxy-carbonyl, hydrophobic or macromolecular carrier gp. attached"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New radio-pharmaceutical agents deriv. from peptide chelating ligands - also the new ligands and complexes with tissue targetting peptide(s), useful for imaging and treatment of e.g.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            DP-178 homologue 36 derived from HIV-1 has antiviral activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  antiviral activity, DP-178; DP-107; diagnostic; HIV-1LAI; human immunodeficiency virus; transmembrane protein; gp41; alpha helix; leucine zipper; DP-185.
                                                                                                                                                                                                                 /note= "Forms peptide linkage with DGlu1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  65.2%; Score 15; DB 15; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Schulze P;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Rohlfs G,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         atherosclerosis or steroid dependent cancer
'note= "D-form residue"
                                                                                                                                           /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Location/Qualifiers
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                                                                  note= "alloDIle"
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/note= "D or L form residues"
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                                                                                                                                                                                                                                                                                                                                                                                     Claim 1; Page 2; 7pp; Japanese.
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les 2; Conserv
                                                                                                                                                                                                                                 (KAOS ) KAO CORP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 AA;
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Modified-site
                    Modified-site
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                                                                             JP06321755-A
                                                                                                                                                      11-MAY-1993;
                                                                                                                                                                                            11-MAY-1993;
                                                                                                                                                                                                                                                                                                                                                 pigmentation
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     11-MAY-1993;
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0
                                                                                                                                                                                                                                                                                                                                                                                                                      truncated at the amino terminus. DP-178 corresponds to amino acids 638 to 673 of the HTV-1 isolate LAI transmembrane protein 1994. It ectodomain, and complexes with DP-107 (corresponds to amino acids 558-595) which contains a leucine zipper motifi The peptides complex activity. The peptide derivs. were identified by a computer assisted peptide sequence search. The peptides inhibit transmission to uninfected cells, and can also be used as type and/or subtype specific diagnostic tools.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                       AAR64365-97 are peptide derivs. of DP-178 (AAR64364) which have been
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "forms peptide bond with Xaa at position 5" /note= "D or L form residues"
                                                                                                                                                                                                                                                                                                          Computer search generated synthetic peptides - are inhibitors of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      í,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           65.2%; Score 15; DB 16; Length 5; llarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Melanin pigmentation inhibition, skin whitening cosmetic, cyclic peptapeptide, sunburn fleck formation treatment.
                                                                                                                                                                                                               Barney SO, Bolognesi DP, Lambert DM, Matthews TJ;
Petteway SR, Wild CT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Melanin pigmentation inhibiting cyclic pentapeptide.
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/note= "D or L form residues"
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/note= "D or L form residues"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'note= "D or L form residues"
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                                                                                                                                                                                                                                                                                                                                                                   Claim 11; Page 133; 182pp; English.
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                                                                                               94WO-US05739.
                                                                                                                                    93US-0073028
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les 2; Conserva
                                                                                                                                                                                                                                                                     WPI; 1995-036105/05
                                                                                                                                                                        (UYDU-) UNIV DUKE.
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                                                                                                                                                                                                                                                                                                                              HIV transmission
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 AA;
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                                                                                             07-JUN-1994;
                                                                                                                                    07-JUN-1993;
                    WO9428920-A
                                                        22-DEC-1994
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Gaps
/note= "forms peptide bond with Leu at position 1"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAR59459-R59461 are cyclic pentapeptides which inhibit melanin pigmentation. As part of a skin whitening cosmetic they can be used for the prevention of fleck formation after sunburn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Skin whitening cosmetic contg. cyclic penta:peptide - e.g. cyclo(L-Leu-D-Trp-D-Glu-L-Ala-D-Val), for inhibiting melanin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Melanin pigmentation inhibition; skin whitening cosmetic; cyclic peptapeptide; sunburn fleck formation treatment.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 5;
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larity 100.0%; Pred. No. 7.8e+05;
Conservative 0; Mismatches 0; Indels
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Gaps

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2
/label= OTHER
/note= "alpha-amino acid with conformationally
/note= "astricted 4-16C side chain, preferably
1,2,3,4-tetrahydro-2-carboline-3-carboxylic
acid"
                                                                                                                                                                                                                                  Endothelin receptor antagonist, organ hypofunction, liver, surgery, transplant, cyclic.
                                                                                                                                                                                                             Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                                                                                                                                                                                                                                                                             /note= "D-Asp, which is part of a cyclic peptide
formed by condensation with D-Trp(5)"
                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "D-Trp, which is part of a cyclic peptide formed by condensation with D-Asp(1)"
                         Length 5;
                                             Indels
                      65.2%; Score 15; DB 16; I
100.0%; Pred. No. 7.8e+05;
iive 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Claim 34; Page 106; 107pp; English.
                                                                                                                                            AAR69429 standard; peptide; 5 AA.
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                                                                                                                                                                                       (first entry)
                                             Conservative
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                               Local Similarity
es 2; Conserv
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   5 AA;
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                                                                                      1 LW 2
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    Sequence
                      Query Match
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                                                                  3:
                                            Matches
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                                                                                                                       RESULT 49
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                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    'note= "forms peptide bond with Asp at position
                                                                                                                                                                                      .;
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                                                                                            AARS9459-R59461 are cyclic pentapeptides which inhibit melanin pigmentation. As part of a skin whitening cosmetic they can be used for the prevention of fleck formation after sunburn.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR59459-R59461 are cyclic pentapeptides which inhibit melanin pigmentation. As part of a skin whitening cosmetic they can be used for the prevention of fleck formation after sunburn.
                              Skin whitening cosmetic contg. cyclic penta:peptide - e.g. cyclo(L-Leu-D-Trp-D-Glu-L-Ala-D-Val), for inhibiting melanin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Skin whitening cosmetic contg. cyclic penta:peptide - e.g. cyclo(L-Leu-D-Trp-D-Glu-L-Ala-D-Val), for inhibiting melanin
                                                                                                                                                                                                                                                                                                                                                                        Melanin pigmentation inhibition, skin whitening cosmetic, cyclic peptapeptide, sunburn fleck formation treatment.
                                                                                                                                                                 Length 5;
                                                                                                                                                             65.2%; Score 15; DB 16; Length 5; ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                    Melanin pigmentation inhibiting cyclic pentapeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                note= "D-form residue"
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                                                                         Example 1; Page 3; 7pp; Japanese.
                                                                                                                                                                                                                                                                                   AAR59461 standard; peptide; 5 AA.
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                                                                                                                                                                          Local Similarity
           WPI; 1995-041168/06.
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                                                                                                                                           5 A.
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Modified-site
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                                                    pigmentation
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                                                                                                                                           Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                           Synthetic.
                                                                                                                                                                                                                             1 LW 2
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                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                    or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ..
Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on transplant of the organ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
65.2%; Score 15; DB 16; Length 5;
Best Local Similarity 100.0%; Pred. No. 7.8e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
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AAR69414 standard; peptide; 5 AA.
AAR69414
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to the new use of known endothelin receptor antagonists for prophylaxis and/or therapy of hypofunction of organs (particularly the liver) which occurs following their surgery or transplant. The present sequence is a specific example of a suitable antagonist listed in the Claims.
                                                                                                                                                                                                            Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                                                                                                                                                                                                       Endothelin receptor antagonist; organ hypofunction; liver; surgery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on or
                                                                                                                                                                                                                                                                                                                                            /note= "D-Ser, which is part of a cyclic peptide
formed by condensation with D-Trp(5)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "D-Trp, which is part of a cyclic peptide
formed by condensation with D-Ser(1)"
                                                                                                                                                                                                                                                                                                                                                                                                     /note= "alpha-amino acid with conformationally restricted side chain"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 65.2%; Score 15; DB 16; Length 5; Best Local Similarity 100.0%; Pred. No. 7.8e+05; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                               Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Kitada C, Wakimasu M, Watanabe T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 34; Page 106; 107pp; English.
                                                                                                                  AAR69430 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                note= "D-Val"
                                                                                                                                                                                                                                                                                                                                                                                        /label= OTHER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     94EP-0106093.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93JP-0094332.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             (TAKE ) TAKEDA CHEM IND LTD.
                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   transplant of the organ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 1995-000740/01.
                                                                                                                                                                                                                                                       transplant; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 AA;
                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                          Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20-APR-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 21-APR-1993;
                                                                                                                                                                              18-JUL-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        30-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP626174-A.
               S
                                                                                                                                                                                                                                                                                    Synthetic.
                                            4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                 AAR69430;
               4 LW
                                                                                    RESULT 50
AAR69430
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/note= "D-Cys(03S), which is part of a cyclic peptide
formed by condensation with D-Trp(5)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The invention relates to the new use of known endothelin receptor antagonists for prophylaxis and/or therapy of hypofunction of organs (particularly the liver) which occurs following their surgery or transplant. The present sequence is a specific example of a suitable antagonist listed in the Claims.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Endothelin receptor antagonist, organ hypofunction, liver, surgery, transplant, cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                                                                                                  Endothelin receptor antagonist; organ hypofunction; liver; surgery;
                                                                                       Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on or transplant of the organ
                                                                                                                                                                                                                                                                                                                                                                                                                    /note= "D-Trp, which is part of a cyclic peptide
formed by condensation with D-Cys(1)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     .;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65.2%; Score 15; DB 16; L. L. L. 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                        Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Kitada C, Wakimasu M, Watanabe T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Claim 12; Page 86; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAR69415 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                         /note= "D-Val"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       94EP-0106093.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    93JP-0094332.
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                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1995-000740/01.
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es 2; Conserv
                                                                                                                                                              transplant; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 AA;
                                                                                                                                                                                                                                                                               Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20-APR-1994;
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                                             18-JUL-1995
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AAR69414;
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The invention relates to the new use of known endothelin receptor antagonists for prophylaxis and/or therapy of hypofunction of organs (particularly the liver) which occurs following their surgery or transplant. The present sequence is a specific example of a suitable antagonist listed in the Claims.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Endothelin receptor antagonist; organ hypofunction; liver; surgery; transplant; cyclic.
                        /note= "D-Trp, which is part of a cyclic peptide formed by condensation with D-Asp(1)"
                                                                                                                                                                                                                                        Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on transplant of the organ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Endothelin antagonist peptide for treating organ hypofunction.
                                                                                                                                                                                                                                                                                                                                                                                                                     65.2%; Score 15; DB 16; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "Tyr-NH-Ind-OH, where Ind is
1-carboxyindane-2-y1"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'note= "hexamethyleneimino-CO-Leu"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                                                          Watanabe T;
                                                                                                                                                                                                                                                                                           Claim 12; Page 86; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAR69392 standard; peptide; 5 AA.
  /note= "D-Val"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      note= "D-Ala"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              'label= D-Trp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               'label= bAla
                                                                                                               94EP-0106093.
                                                                                                                                        93JP-0094332
                                                                                                                                                                 (TAKE ) TAKEDA CHEM IND LTD
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Best Local Similarity 100.v.
Best Zi Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             16-JUL-1995 (first entry)
                                                                                                                                                                                         Wakimasu M,
                                                                                                                                                                                                                 WPI; 1995-000740/01.
              Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                             5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
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                                                                                                              20-APR-1994;
                                                                                                                                        21-APR-1993;
                                                                                       30-NOV-1994.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    30-NOV-1994
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                                                              EP626174-A
                                                                                                                                                                                         Kitada C,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR69392;
                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 54
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                                                                                                                                                                                                                                                                                                                                                                                                                                                     The invention relates to the new use of known endothelin receptor antagonists for prophylaxis and/or therapy of hypofunction of organs (particularly the liver) which occurs following their surgery or transplant. The present sequence is a specific example of a suitable antagonist listed in the Claims.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Endothelin receptor antagonist; organ hypofunction; liver; surgery;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cyclic endothelin receptor peptide for treating organ hypofunction.
                                                                                                       /note= "D-Thg(2) or D-Cpg, where Thg(2) is 2-(2-thienyl)glycine and Cpg is 2-cyclopentyl-glycine"
                                                                                                                                                                                                                                                                                                                                                                            Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on or transplant of the organ
                                                            /note= "D-Asp, which is part of a cyclic peptide
formed by condensation with D-Trp(5)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "D-Asp, which is part of a cyclic peptide
formed by condensation with D-Trp(5)"
                                                                                                                                                             /note= "D-Trp, which is part of a cyclic peptide
formed by condensation with D-Asp(1)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
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                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
                                                                                                                                                                                                                                                                                                                              Watanabe T;
                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 12; Page 86; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAR69413 standard; peptide; 5 AA.
                                                                                                 /label= OTHER
                                                                                                                                                                                                                                                    94EP-0106093
                                                                                                                                                                                                                                                                             93JP-0094332
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                18-JUL-1995 (first entry)
                                                                                                                                                                                                                                                                                                                            Kitada C, Wakimasu M,
                                                                                                                                                                                                                                                                                                                                                      WPI; 1995-000740/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                transplant; cyclic
                                                                                                                                                Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 AA;
                                                                                   Misc-difference
                                               Misc-difference
                                                                                                                                                                                                                                                  20-APR-1994;
                                                                                                                                                                                                                                                                           21-APR-1993;
                                                                                                                                                                                                                         30-NOV-1994
                                                                                                                                                                                                   EP626174-A
           Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Seguence
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Synthetic.
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                                                                                                                                                     The invention relates to the new use of known endothelin receptor antagonists for prophylaxis and/or therapy of hypofunction of organs (particularly the liver) which occurs following their surgery or transplant. The present sequence is a specific example of a suitable antagonist listed in the Claims.
                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       part of a cyclic structure obtained by condensation of D-Trp(5) onto the alpha-NH2"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   part of a cyclic structure formed by condensation of the COOH onto the NH2 group of D-Glu(1)"
                                                                                                                                                                                                                                                                                                                                                                                                     diagnostic imaging, atherosclerotic plaque; tumour; inflammation; conjugate; radiotherapy; cysteine-free; metal-binding; endothelin;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "D-form residue; and this amino acid forms
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "D-form residue; and this amino acid forms
                                                                                                Use of endothelin receptor antagonists - for prophylaxis or treatment of organ hypofunction, partic. caused by surgery on or
                                                                                                                                                                                                                                                                                                                                                                                   Endothelin sequence for use as conjugate with Cys-free peptide.
                                                                                                                                                                                                                                               ;
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                                                                                                                                                                                                                             65.2%; Score 15; DB 16; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                               Kitada C, Wakimasu M, Watanabe T;
                                                                                                                                    Claim 7; Page 83; 107pp; English.
                                                                                                                                                                                                                                                                                                                               AAR86453 standard; peptide; 5 AA.
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          94EP-0106093.
                                            (TAKE ) TAKEDA CHEM IND LTD.
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                           93JP-0094332
                                                                                                                                                                                                                                                                                                                                                                   05-FEB-1996 (first entry)
                                                                                                                                                                                                                                               Conservative
                                                                                                                    transplant of the organ
                                                                                                                                                                                                                                       Local Similarity
ses 2; Conserva
                                                                               WPI; 1995-000740/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference 5
                                                                                                                                                                                                            5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
         20-APR-1994;
                           21-APR-1993;
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Synthetic.
                                                                                                                                                                                                                                                                 4 LW 5
                                                                                                                                                                                                                                                                                    1 LW 2
                                                                                                                                                                                                                                                                                                                                                 AAR86453;
                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                       cyclic.
                                                                                                                                                                                                                                                Matches
                                                                                                                                                                                                                                                                                                              RESULT 55
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New peptides of formula R1-X-R2 are provided, together with their conjugates with peptides, proteins, biomolecules and macromolecules, their complexes with metal ions and their water-soluble salts; where their complexes with metal ions and their water-soluble salts; where including at least one Met, Arg, Lys or Asn but no Cys; R1 is H or a hydrocarbyloxy or an optionally substituted amine group (forming an hydrocarbyloxy or an optionally substituted amine group (forming an metal (e.g. 99m-T0) for in-vivo diagnostic imaging and therapy of tumours, organs and foci of inflammation. Conjugates of the peptides can be targetted to specific itssues. A particular application is imaging of the prosclerotic plaque. Metal is bound with good in-vivo stability; unbound material is rapidly cleared; and there is no allergic reaction. Specific examples of the peptides are paptides are free of Cys, they can be prepared simply and rapidly and can be presented in the form of a kit for radioparameterial is rapidly. Specific examples of the peptides are given in ARR8444 - ARR84439.

The present sequence is a specific example of an endothelin peptide which can be used as a conjugate with the new peptides.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                sex-hormone, human; veterinary medicine; cancer; prostate; uterus; breast; pituitary; prostatomegaly; endometriosis; hysteromyoma; precocious puberty; amenorrhoea; premenstrual syndrome; oylic; multilocular ovarian syndrome; comedo; pregnancy; contraception; ovulation; menstrual cycle; cestrus; meat quality; growth; spawning; fish; testosterone; superagonist; leuprorelin acetate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "N-pToluenesulphonylarginine, D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Luteinising hormone releasing hormone receptor; LH-RH; antagonist;
        Frommel C, Hohne W;
ier HM, Schneider-Mergener J;
    Conrad J, Dinkelborg L, Erber S, Frommel C, Hohne W;
Kramp W, Kuttner G, Malin R, Schier HM, Schneider-Mergener Steinbrecher R, Froemmel C, Hoehne W, Kuettner G, Schier H;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        'note= "Joined via a peptide linkage to Trp5"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "Joined via a peptide linkage to Phel"
                                                                                                                                                                                                                                       New cysteine-free peptide(s) and their metal ion complexes conjugated to targetting agent, useful for in vivo imaging tumours, atherosclerotic plaque etc. and for radiotherapy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cyclic pentapeptide #25 used in LH-RH receptor antagonist.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  65.2%; Score 15; DB 16; L
llarity 100.0%; Pred. No. 7.8e+05;
Conservative 0; Mismatches 0;
Erber S, Frommerin R. Schier HM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAW17809 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                               Claim 13; Page 20; 25pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07-JUL-1997 (first entry)
                                                                                                                                                                   WPI; 1995-171400/23.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW17809;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Peptide mimetic; endothelial leukocyte adhesion molecule; ELAM; selectin; receptor; leukocyte; vascular wall; endothelium; extravasation; inflammation; sialyl Lewis; cell surface glycoprotein; HL60 cell.
                                                                                                                                                                                                                                      intentialing hormone releasing hormone (LHT.RH) receptor antagonist composition of the invention. Peptides such as this are used to prevent or treat sex-hormone dependent disorders in human or veterinary medicine, esp. cancers (of the prostate, uterus, breast or pituitary), prostatomegaly, endometriosis, hysteromyoma or precocious puberty, but also amenorrhoes, premenstrual syndrome, multilocular ovarian syndrome, comedo, etc.; also to control pregnancy (contraception for men or women, also to induce ovulation) and the menstrual cycle. They are also used growth, and to promote spawning in fish. They may also inhibit the translent increase in testosterone blood levels caused by admin. of superagonists such as leuprorelin acetate.
                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                This peptide represents a cyclic peptide which is included in the
                                                                                                                                                           Cyclic penta:peptide(s), some new, as LH-RH receptor antagonists used to treat or prevent sex hormone dependent disorders, e.g. cancer, also for control of pregnancy and menstruation and to improve meat quality in animals
                                                                                                                                                                                                                                                                                                                                                                                                                  .;
0
                                                                                                                                                                                                                                                                                                                                                                                           Score 15; DB 17; Length 5;
Pred. No. 7.8e+05;
0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "amidated C-terminus"
                                                                                                                                                                                                           Example 45; Page 173; 198pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anti-ELAM-1 binding peptide #213.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR86236 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   'note= "Ac-Leu"
                                                                                                                     Furuya S, Kato K, Kitada C;
                                                                                                                                                                                                                                                                                                                                                                                           65.2%;
                                                                   95JP-0110933.
                                                  96WO-JP01140
                                                                                                 (TAKE ) TAKEDA CHEM IND LTD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             27-JUN-1996 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                2; Conservative
                                                                                                                                        WPI; 1996-497569/49.
                                                                                                                                                                                                                                                                                                                                                                                                     Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Key
Modified-site
                                                25-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Modified-site
                                                                   09-MAY-1995;
                                                                              28-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                  1 FXXLW 5
                              31-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WO9531210-A1
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                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR86236;
                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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that bind to endothelial leukocyte adhesion molecule (BLAM)-1. This that bind to endothelial leukocyte adhesion molecule (BLAM)-1. This molecule is a member of the selectin family of receptors and is involved in binding of leukocytes to the vascular endothelial wall prior to extravasation of the leukocyte, e.g. to a site of inflammation.

The peptides bind pref to B-selectin but may also bind L- or P-selectin, and can be used to treat conditions mediated by E-selectin, e.g. inflammatory conditions. The peptides have strong affinity for the selectin receptors and inhibit the binding of the sialyl Lewis (SLE-x) car of cell surface glycoproteins to B-selectin. The peptide are small, generally less than 2 kD, have an IC50 of up to 100 micromole against binding of HL60 cells to BLAM-1, have one or more peptide linkages replaced by CH2OC(O)NR, phosphonate, CH2SOZNR, CH3ONR, CON(R6), con NHCONH linkages where R = H or a lower alkyl and R6 = a lower alkyl. The peptides may also have substituted N- and C-termini e.g. cucinimido, N-benzyloxycarbonyl or N-lower alkyl cpds.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                 New peptide(s) that bind to endothelial leukocyte adhesion molecule 1 - useful for treating inflammation and other E-selectin mediated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Bifunctional, bis-nicotinamide, chelator, radioactive, technetium, rhenium, conjugate, endothelin, angiotensin, targetting agent, organ; in vivo imaging, receptor; tissue, atherosclerotic plaque; diagnosis, breast, prostatic, carcinoma; cytostatic agent;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Targetting conjugate for bis-nicotinamide bifunctional chelator.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Indels
                                                                                                                                                                                                Barrett RW, Cwirla SE, Dower WJ, Koller KJ, Lee J;
Martens CL, Ruhland-fritsch B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 17; L 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          note= "allo-D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note= "D-form residue"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW33328 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 3; Page 73; 85pp; English.
                                                                                                                                  (AFFY-) AFFYMAX TECHNOLOGIES NV
                                                                        94US-0241054.
      95WO-US06315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         05-FEB-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                   WPI; 1996-010687/01.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Misc-difference 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 delivery; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference
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11-MAY-1995;
                                                                 11-MAY-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW33328;
                                                                                                                                                                                                                                                                                                                                                                                                                              diseases
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW33328
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Matsuo Y, Ninomiya M;
                                                                                                     WPI; 1997-034102/03.
                                                                                                                                                                                                                                                                                                             Local Similarity
les 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  delivery; cyclic
                                                                                                                                                                                                                                                                                    5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DE19536785-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Homo sapiens.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  21-SEP-1995;
                     29-MAY-1996;
                                        02-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          27-MAR-1997
05-DEC-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW33020;
                                                                                                                                                                                                                                                                                    Sequence
                                                                                                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                                                                    4 LW
                                                                                                                                                                                                                                                                                                                             Matches
                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 60
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The alpha carboxyl group is condensed with the alpha amino group of Asp (1)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        /note= "D-form residue. This Asp is not a true N-terminal residue but is part of a cyclic peptide.
The alpha amino group is condensed with the alpha carboxyl group of Trp(5)."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       cerebral hernia; tumour; apoplexy; hypertension; respiratory failure; hyperglycaemia; hypoglycaemia; adrenal function failure; collagen disease; poisoning; cyclic.
                                                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                    Novel bis-nicotinamide bifunctional chelating agent for radioactive technetium and rhennium, when conjugated to an endothelin or angiocensin derived targetting agent, e.g. the present peptide, can be used for the in vivo imaging of organs, receptors and receptor containing tissues and/or atherosclerotic plaques, e.g. for the diagnosis of breast or prostatic carcinoma. They can also be used for cytostatic agent delivery.
                                                                                                                                                         New bis-nicotinamide bi:functional chelating agents - forming new, stable technetium or rhenium complexes and conjugates useful as radiodiagnostic and radio:therapeutic agents
                                                                                                                                                                                                                                                                                                                                                  .;
0
                                                                                                                                                                                                                                                                                                                             Length 5;
                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Endothelin antagonist #1 used as brain oedema inhibitor.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Endothelin antagonist; brain oedema inhibitor; bosentan;
                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 18; L Llarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0;
                                                                                                       Hilger CS, Kramp W, Platzek J;
                                                                                   (UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW32229 standard; peptide; 5 AA.
                                                                                                                                                                                                    Claim 11; Page 42; 47pp; German.
                                          96WO-DE01824.
                                                              95DE-4036783.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-FEB-1998 (first entry)
                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
                                                                                                                                      WPI; 1997-212562/19.
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                                                                                                                                                                                                                                                                                                          5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Modified-site
                                                                                                       Dinkelborg L,
Raduechel B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Modified-site
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  WO9710853-A2
                                           19-SEP-1996;
                                                               21-SEP-1995;
                       27-MAR-1997.
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                                                                                                                                                                                                                                                                                                           Sequence
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This peptide is an endothelin antagonist which is the active component in a new cerebral oedema inhibitor, used to prevent swelling in head injury, cerebral tumour, blood vessel damage in apoplexy, hypertension, respiratory failure, carbon monoxide poisoning, hyponatraemia, acute Kidney damage, hyperglycaemia, hypoglycaemia, adrenal function failure, collagen disease, inadequate buffering of blood, or poisoning by tin, lead or arsenic. The inhibitor prevents cerebral hernia, headache, nausea, dizziness, restlessness and loss of consciousness, especially sudden death from cerebral hernia.
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                                                                                                                                                                                                                                                                                                                                                                               Cerebral oedema inhibitor comprising endothelin antagonist - esp. specified oleanonic acid deriv., cyclopeptide or bosentan
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llarity 100.0%; Pred. No. 7.8e+05;
Conservative 0; Mismatches 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 2; Page 12; 20pp; Japanese.
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96WO-JP01438.
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Homo sapiens.
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                                                                                                                                                               Bifunctional sulphonamide type ligands for radioactive technetium or rhenium - and their chelates and conjugates, useful for imaging
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                     technetium and rhenium, when conjugated to an endothelin or angiotensin derived targetting agent, e.g. the present peptide, can be used for the in vivo imaging of organs, receptors and receptor containing tissues and/or atherosclerotic plaques, e.g. for the diagnosis of breast or prostatic carcinoma. They can also be used for cytostatic agent delivery.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       rhenium, conjugate, endothelin, angiotensin, targetting agent, organ; in vivo imaging; receptor; tissue, atherosclerotic plaque; diagnosis; breast; prostatic; carcinoma; cytostatic agent;
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                                                                                                                                                                                                                                                                                                               Novel bifunctional sulphonamide type ligands for radioactive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bifunctional sulphonamide; ligand; radioactive; technetium;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 18; Length 5; 100.0%; Pred. No. 7.8e+05; ative 0; Mismatches 0; Indels
                                                 Platzek J;
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                                                 Kramp W,
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(UYBE-) UNIV BERLIN INST DIAGNOSTIKFORSCHUNG.
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                                                 Hilger CS,
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                                                                                                                                                                                                                                                                  Claim 9; Page 12; 12pp; German.
                                                                                                                                                                                                                carcinoma or atherosclerosis
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                                                                                                                    WPI; 1997-193894/18.
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                                               Dinkelborg L,
Raduechel B;
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Raduechel B;
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Bifunctional sulphonamide type ligands for radioactive technetium and rhenium - and their chelates and conjugates with targetting agents, useful for imaging carcinoma, atherosclerotic plague, etc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Bifunctional sulphonamide, ligand, radioactive, technetium, thenium, conjugate, endothelin, angiotensin, targetting agent, organ; in vivo imaging; receptor; tissue, atherosclerotic plaque; diagnosis, breast, prostatic, carcinoma; cytostatic agent,
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                                                                                                    Novel bifunctional sulphonamide type ligands for radioactive
                                                                                                                                                                                                                                                                            Score 15; DB 18; Length 5;
Pred. No. 7.8e+05;
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100.0%; Pred. No. ...
0; Mismatches
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                                                                     Claim 11; Page 19; 19pp; German
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                                                                                                                                                                                                                                                                                              Local Similarity
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Raduechel B;
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          Novel bifunctional sulphonamide type ligands for radioactive technetium and rhenium, when conjugated to an endothelin or angiotensin derived targetting agent, e.g. the present peptide, can be used for the in vivo imaging of organs, receptors and receptor containing tissues and/or atherosolerotic plaques, e.g. for the diagnosis of breast or prostatic carcinoma. They can also be used
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                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conjugating cyclic peptide for radio-therapeutic/diagnostic agent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conjugate; diseased tissue; endothelin; angiotensin; chemotactic; radiodiagnostic; radiotherapeutic; diagnosis; tumour; ischaemia; atherosclerosis; vascular disorder; ischemia; circular; cyclic.
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                                                                                                                                                                           Length 5;
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                                                                                                                                                                         65.2%; Score 15; DB 18; L
100.0%; Pred. No. 7.8e+05;
tive 0; Mismatches 0;
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                                                                                                             for cytostatic agent delivery.
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                                                                                                                                                                                                          Conservative
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                                                                                                                                                            Query Match
Best Local Similarity
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                                                                                                                                          5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide binding to low density lipoprotein - useful in removing LDL from body fluids and as carrier peptide(s) for drugs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present low density lipoprotein (LDL) binding peptide can be used as an absorbent to remove LDL from body fluids, or as a LDL binding reagent. The peptide can be prepared easily and at low cost, and has excellent stability and preservability. It specifically binds LDL, and does not cause bradykinin production, white blood cell activation or blood agglutination.
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                                                                                                                                                                                                                                                                                        density; lipoprotein; LDL; binding; absorbent; removal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 18; Length 5; 100.0%; Pred. No. 7.8e+05; tive 0; Mismatches 0; Indels
                                                 Length 5;
                                                                         0; Indels
                                                65.2%; Score 15; DB 18; L
100.0%; Pred. No. 7.8e+05;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                Low density lipoprotein binding peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Example 11; Page 50; 61pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                      Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                               /note= "amidated"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAW22920 standard; peptide; 5 AA.
                                                                                                                                                                                       AAW22915 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (ASAH ) ASAHI KASEI KOGYO KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (ASAH ) ASAHI MEDICAL CO LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                95JP-0176904.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        96WO-JP01734.
                                                                                                                                                                                                                                          02-OCT-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Aritomi M, Hatanaka Y;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
nes 2; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-087320/08.
                                                                                                                                                                                                                                                                                                      body fluid; reagent
vascular disorders.
                                                         Local Similarity
ses 2; Conserv
                          5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 AA;
                                                                                                                                                                                                                                                                                                                                                                  Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                      WO9700889-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                        21-JUN-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                09-JAN-1997
                                                                                                   I.W.
                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                           4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        4 LW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                'n
                                                                                                                                                                                                                AAW22915;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Seguence
                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                4 LW
                                                                                                    4
                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
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                                                                                                                                                                RESULT 64
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ID AAW2
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WO9700889-A1
                                                                                                                                                                                                                                                                                         Matches
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AAY20414
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                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                      Peptide binding to low density lipoprotein - useful in removing LDL from body fluids and as carrier peptide(s) for drugs
                                                                                                                                                                                                                                                                                                                                    The present low density lipoprotein (LDL) binding peptide can be used as an absorbent to remove LDL from body fluids, or as a LDL binding reagent. The peptide can be prepared easily and at low cost, and has excellent stability and preservability. It specifically binds LDL, and does not cause bradykinin production, white blood cell activation or blood agglutination.
                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
                                                              Low density, lipoprotein; LDL; binding; absorbent; removal; body fluid; reagent.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Low density; lipoprotein; LDL; binding; absorbent; removal; body fluid; reagent.
                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 18; Length 5; 100.0%; Pred. No. 7.8e+05; rative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Low density lipoprotein binding peptide.
                                           Low density lipoprotein binding peptide
                                                                                                            Location/Qualifiers
                                                                                                                                                                                                                                                                                                                  Example 11; Page 50; 61pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                     5
/note= "amidated"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAW22928 standard; peptide; 5 AA
                                                                                                                                                                                                                      (ASAH ) ASAHI KASEI KOGYO KK.
(ASAH ) ASAHI MEDICAL CO LTD.
                                                                                                                                                                                                      95JP-0176904.
                                                                                                                                                                                     96WO-JP01734.
                          02-OCT-1997 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              02-OCT-1997 (first entry
                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                   Aritomi M, Hatanaka Y;
                                                                                                                                                                                                                                                                      WPI; 1997-087320/08.
                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                      5 AA;
                                                                                                                    Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Key
Modified-site
                                                                                                                                                                                    21-JUN-1996;
                                                                                                                                                                                                     21-JUN-1995;
                                                                                                                                                WO9700889-A1
                                                                                                                                                                 09-JAN-1997
                                                                                           Synthetic
         AAW22920;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAW22928;
                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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LV
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Human; beta-amyloid precursor protein; beta-APP; diagnosis; cancer; frameshift mutation; age-related disease; neurodegenerative disorder; Alzehmer, s disease; Down's syndrome; myotonic dystrophy; neuronal; Huntington's disease; multiple sclerosis; alcoholic liver disease; diabetes mellitus type II; microtubule associated protein; Tau; Big Tau; ubiquitin B; apolipoprotein E; MAP2; neurofilament-L; neurofilament-C; presenilin I; presenilin II; cellular tumour antigen; glial fibrillary acidic protein; GFAP; p53; semaphorin III; HUPF-I; bcl-2; B-cell leukemia/lymphoma 2 proto-oncogene; HMGP-C; MSP-A; high mobility group protein-C; neuroendocrine specific protein A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                    Peptide binding to low density lipoprotein - useful in removing LDL from body fluids and as carrier peptide(s) for drugs
                                                                                                                                                                                                                                                                                                                                                                      The present low density lipoprotein (LDL) binding peptide can be used as an absorbent to remove LDL from body fluids, or as a LDL binding reagent. The peptide can be prepared easily and at low cost, and has excellent stability and preservability. It specifically binds LDL, and does not cause bradykinin production, white blood cell activation or blood agglutination.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65.2%; Score 15; DB 18; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
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(ROYA-) ROYAL NETHERLANDS ACAD ARTS & SCI.
(UYRO-) UNIV ROTTERDAM ERASMUS.
                                                                                                                                                                                                                                                                                                                                     Example 11; Page 50; 61pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAY20414 standard; Protein; 5 AA.
                                                                                                                       (ASAH ) ASAHI KASEI KOGYO KK.
(ASAH ) ASAHI MEDICAL CO LTD.
                                                                                  95JP-0176904.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            97US-0043163.
                                         96WO-JP01734
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                        Aritomi M, Hatanaka Y;
                                                                                                                                                                                                                             WPI; 1997-087320/08
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 AA;
                                         21-JUN-1996;
                                                                                  21-JUN-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              10-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
09-JAN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           15-OCT-1998.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAY20414;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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proteins Tau and Big Tau, ubiquitin B, apolipoprotein E, microtubule associated protein 2 (MAP2), neurofilament-L, neurofilament-M, neurofilament-F, presentlin I, glala fibrillary acidic protein (GPAP), the cellular timour antigen p53, B-cell leukenia/lymphoma 2 (bcl.2) prote-oncogene, semaphorin III, HUPF-1, high mobility group protein-C (HMGP-C) and neuroendocrine specific protein A.
                                                                                                                                                                                                                                                 This invention describes a novel method for the diagnosis of a disease caused by, or associated with, an RNA molecule that has a frameshift mutation. The method is used to diagnose age-related diseases, especially cancer and a wide range of neurodegenrative disorders (e.g. Alzheimer's disease, Down's syndrome, myotonic dystrophy, Huntington's disease, and sease, nearlitus type II and many others listed) or susceptibility to these disorders. The method allows a definitive diagnosis of Alzheimer's disease in living patients, at an early stage. It is based on the observation that disease may be caused by mutations in RNA rather than DNA. The invention describes the used of neuronal system RNA molecules, specifically proteins including beta-amyloid precursor protein (beta-APP), the microtubule associated
                                                                                                    Diagnosing disease by detecting frameshift mutations in RNA or corresponding protein mutations - used to diagnose cancer and neurological diseases, particularly Alzheimer's disease, and also for treatment and prevention with specific ribozymes or wild-type
                 Van Leeuwen FW;
                                                                                                                                                                                                                   Disclosure; Figure 6; 258pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW65861 standard; peptide; 5 AA.
                 Burbach JPH, Grosveld FG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-OCT-1998 (first entry)
                                                   WPI; 1998-609901/51.
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es 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            5 AA;
                                                                      N-PSDB; AAX75757
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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Matches
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%XTXXBXXFFFFFFFXX8XXCCCCCCCXX8XXFFFFFFFFFFXX8XXFFX
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0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "D-Glu is condensed onto D-Trp at position 5 to form a cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /note= "D-Trp is condensed onto D-Glu at position 1 to form a cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           tyrosine kinase blocker; chemotherapy; antibody; conjugate; cyclic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              endothelin receptor; antagonist; atherosclerosis; radionuclide;
65.2%; Score 15; DB 19; Length 5; llarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Endothelin receptor binding peptide part of conjugate.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /label= alle
/note= "D-form allo-isoleucine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
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0; Gaps

Indels

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Cyclic peptide which mimics surface feature of endothelin.

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31-JUL-1998 (first entry)

AAW50857;

AAW50857 standard; peptide; 5 AA.

RESULT 69 AAW50857 Endothelin; endothelina; endothelinb; ETa; ETb; receptor; endothelin-1; cyclic.

Synthetic.

10-JUN-1998,

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The invention relates to (A) the use of conjugates of the following formula as therapeutic agents: B-Wn, where B = a residue which can bind an endothelin receptor, and is derived from an endothelin partial sequence or an endothelin antagonist; W = an active group which: (i) is a radionuclide or (ii) is derived from a chemotherapeutic agent, a complex with a cardiovacte, a noligonucleotide, a protein tyrosine kinase complex a carbohydrate, an oligonucleotide, a protein tyrosine kinase blocker, an anti-thrombotic agent, a growth factor inhibitor, an anti-thrombotic agent, a growth factor inhibitor, an anti-thrombotic agent, a growth factor inhibitor, an anti-thrombotic agent, a scowth factor inhibitor, an anti-thrombotic agent, a scowth factor inhibitor, an anti-thrombotic agent, a lipid lowering agent or an anti-thrombotic agent, a scowth factor inhibitor, an anti-thrombotic agent, a scowth as a conjugates become enriched in cells in which endothelin receptors conjugates become enriched in cells in which endothelin receptor binding conjugates become enriched sites can be achieved. Unbound conjugate conjugates a specifically craimed endothelin receptor binding conj
                                                                                                                                                                                                                                                                         Therapeutic use, e.g. in treatment of atherosclerosis, of endothelin conjugates - which comprise residue which can bind endothelin receptor, conjugated to groups such as radionuclides or protein tyrosine kinase blockers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65.2%; Score 15; DB 19; L ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0;
                                                                                                                                                                             Dinkelborg L, Hilger C,
                                                                                                                                                                                                                                                                                                                                                                                                                   Claim 3; Page 15; 21pp; German.
                      96DE-1052374.
                                                                     96DE-1052374.
                                                                                                                         SCHD ) SCHERING AG.
                                                                                                                                                                                                                            WPI; 1998-363559/32.
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Matches 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 AA;
                      04-DEC-1996;
                                                                          04-DEC-1996;
                                                                                                                                                                             Blume F,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
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/note=
                                                                                                                                                                                                                    Chan MF;
                                                                                                                                                                                                                                      WPI; 1998-239252/21.
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Best Local Similarity
                          Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-OCT-1999
                                             Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        12-NOV-1996;
                                                                                                                                10-NOV-1994;
                                                                                                                                                   10-NOV-1994;
                                                                                                                                                          14-DEC-1990;
18-JUN-1992;
                                                                                                                                                                               05-APR-1994;
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                                                                                            US5736509-A
                                                                                                              07-APR-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                           4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAY30995
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ô
                                                                                                                                                                                                                                                                                                                                          The invention relates to cyclic peptides that mimic the surface features of the endothelin peptides. The peptides shown in the invention are pentapeptides, hexapeptides and heptapeptides that contain an L-Trp, D-Trp residue or a derivative of D- or L-Trp, such as N-Me-Trp. The peptides modulate the activity of one or more members of the endothelin family of pepides. They are useful for relieving symptoms of and for treating endothelin mediated disorders and diseases. The present sequence represents a cyclic pentapeptide.
                                                                                  /note= "D-form residue, carboxy group of Trp is
condensed onto the amino group of Glu1 to form
a cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                   New cyclic penta-peptides which are mimics of C-terminus of natural endothelin-1 - useful for relieving symptoms of and for treating endothelin mediated disorders and diseases
                          /note= "D-form residue, amino group of Glu is
condensed onto the carboxy group of Trp5 to
form a cyclic peptide"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cyclic peptide which mimics surface feature of endothelin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                  65.2%; Score 15; DB 19; Length 5; 100.0%; Pred. No. 7.8e+05; rative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Endothelin; endothelina; endothelinb; ETa; ETb; receptor;
                                                                  /note= "D-form residue'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
1
                                                                                                                                                                                                                                                                                                                           Example 3; Column 33; 30pp; English
          Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAW50859 standard; peptide; 5 AA.
                                                                                                                                                                                                                               (TEXA-) TEXAS BIOTECHNOLOGY CORP.
                                                                                                                                                                               94US-0410955.
90US-0628111.
92US-0900623.
94US-0223513.
                                                                                                                                                              94US-0410955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              31-JUL-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           endothelin-1; cyclic.
                                                                                                                                                                                                                                                  Chan MF;
                                                                                                                                                                                                                                                                  WPI; 1998-239252/21.
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                        Misc-difference 3
                                                                                                                                                                                                                                                                                                                                                                                                                                 5 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Key
Modified-site
                   Modified-site
                                                                          Modified-site
                                                                                                                                                              10-NOV-1994;
                                                                                                                                                                                                            05-APR-1994;
                                                                                                                                                                                 10-NOV-1994;
                                                                                                                                                                                         14-DEC-1990;
                                                                                                                                                                                                   18-JUN-1992;
                                                                                                                          US5736509-A
                                                                                                                                          07-APR-1998
                                                                                                                                                                                                                                                  Balaji VN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic
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Matches
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The invention relates to cyclic peptides that mimic the surface features of the endothelin peptides. The peptides shown in the invention are pentapptides, hexapeptides and heptiapptides that contain an L-Trp, D-Trp residue or a derivative of D- or L-Trp, such as N-Me-Trp. The peptides modulate the activity of one or more members of the endothelin family of pepides. They are useful for relieving symptoms of and for treating endothelin mediated disorders and diseases. The present sequence represents a cyclic pentapeptide.
                                                                                                                                                                                                "D-form residue, carboxy group of Trp is condensed onto the amino group of Aspl to form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New cyclic penta-peptides which are mimics of C-terminus of natural endothelin-1 - useful for relieving symptoms of and for treating endothelin mediated disorders and diseases
/note= "D-form residue, amino group of Asp is
condensed onto the carboxy group of Trp5
form a cyclic peptide"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Pred. ...
                                                                                                                                                                                                                                                              a cyclic peptide"
                                                                                                                                  'note= "D-form residue"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Example 5; Column 33; 30pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (TEXA-) TEXAS BIOTECHNOLOGY CORP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAY30995 standard; peptide; 5 AA.
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This invention describes a novel aqueous suspension of monodisperse particles on non-crosslinked, non-denatured albumin (50-5000 mm) which is stable against dissolving upon dilution with an alcohol free aqueous medium. The method involves (a) forming an aqueous solution containing albumin and hemoglobin and (b) treating the aqueous solution with an alcohol to cause the solution to become turbid. The particles are useful as agents for in vivo administration, either of their own administration or as a vehicle for other therapeutic or diagnostic agents. The method permits the formation of albumin and hemoglobin particles in the nanometer and micrometer size range, in a form closer to their natural form than the forms of the prior art. The particles therefore constitute a more closely controlled agent for in vivo administration, with greater case of clearance from the body after their period of usefulness.

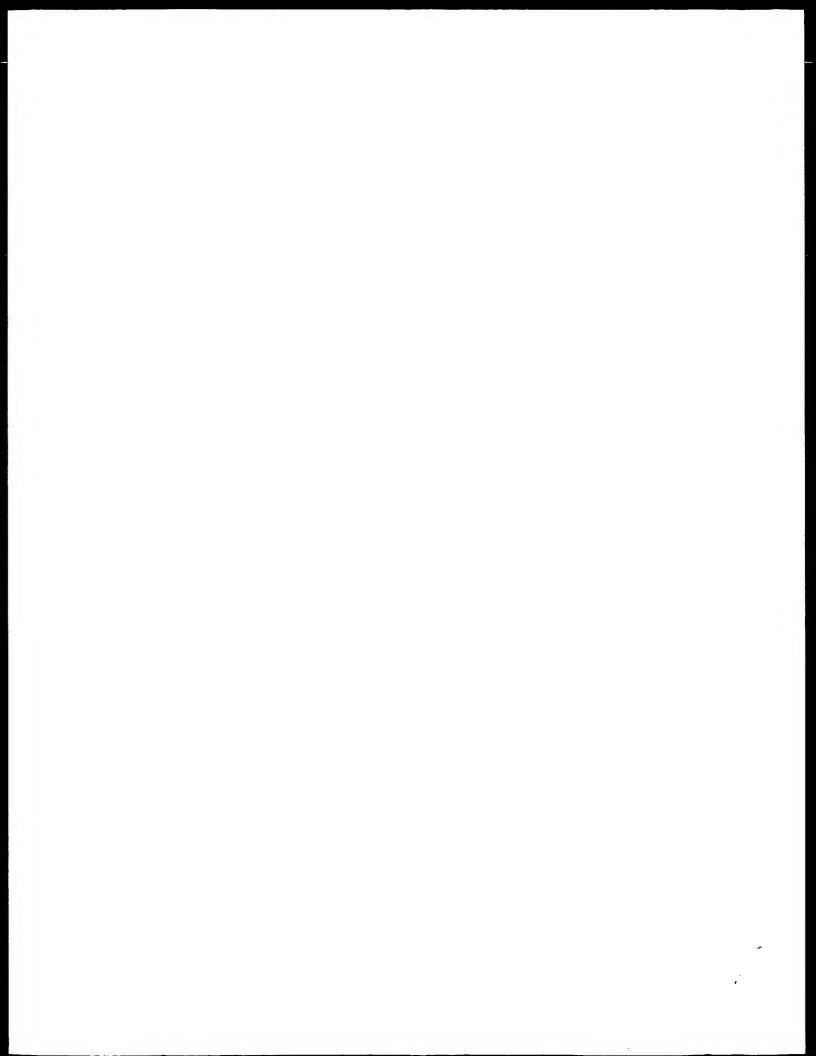
AAY30952-Y31135 represent peptides used in the method of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gene characterisation; recombinase; homology clamp; homology motif tag; HMT; genetic manipulation; genetic engineering; gene transcripion; GPCR; drug target; homologous recombination; target identification; TGF-beta; G-protein coupled receptor; transforming growth factor-beta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
                                                                                                                                                                                                            Non-crosslinked protein particles for therapeutic and diagnostic use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Acetyl choline (muscarinic) GPCR consensus peptide motif.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 20; L ilarity 100.0%; Pred. No. 7.8e+05; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                           Example 22; Column 65-66; 65pp; English.
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               94US-0212546.
91US-0641720.
92US-0959560.
                                                               93US-0069831.
96US-0747137.
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                                                                                                              (HEMO-) HEMOSPHERE INC.
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Best Local Similarity
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                                                                                                                                                                            WPI; 1999-508153/42.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         5 AA;
                                                                             12-NOV-1996;
               14-MAR-1994;
15-JAN-1991;
                                              13-OCT-1992;
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                                                               01-JUN-1993;
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                                                                                                                                             Yen RCK;
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The invention provides compositions and methods for the evaluation and characterisation of individual and sets of genes in disease states. The composition comprises at least one recombinase and at least two single-cranded targeting polynucleotides which are substantially complementary to each other and each having a consensus homology clamp for a gene family i.e. a homology motif tag (HMT). The composition is useful in kit form which include the composition as libraries or pools of degenerate cssDNA probes along with other reagents such as recombinase etc. The methods and compositions are used for inactivation of a gene family gene concerns and compositions are used for inactivation of a gene family gene corresponds and compositions are used for inactivation of a gene family gene corresponds and compositions are used for inactivity of one or more genes in a cell control of disease, or in the elucidation of gene function and activity. Alternatively, the biological activity of the wild-type gene carivity. Alternatively, the biological activity of the wild-type gene carivity. Alternatively, the biological activity of the mind-type gene sequences that affect the transcription of genes, including promoter, repressors, enhancers and transcriptional activating sequences. The compositions are useful in functional activating sequences. The compositions are useful in functional activating sequences. The compositions are useful in functional activating sequences. The composition of new drug targets. HMTs used in homologous recombination compositions are useful in functional genemate studies as well as in identification of new drug targets. HMTs used in homologous recombination and callular level to identify genes involved in a cellular level to identify genes involved in a cellular level to identify genes including those related to disease states. This including those related to disease states. This may also be consumed to the proper description of the proper description of the proper description of the proper description
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                phenotypes i.e. target identification. Sequences AAY31248-308 represent consensus peptide motifs of the G-protein coupled receptor (GPCR) gene family and transforming growth factor-beta (TGF-beta) gene family.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
                       New compositions and methods for targeting sequence modifications in related family genes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Antigenic peptide, streptokinase, streptokinase-specific antibody, thrombolytic activity, thrombolytic therapy, glomerulonephritis,
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                                                                                                Disclosure, Fig 1A; 46pp; English.
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Matches 2; Conserva
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Job time : 82 secs
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                                                                                                                                                                                                                                                                                        The present sequence represents an antigenic peptide of streptokinase. The specification describes a polypeptide which binds to a streptokinase-specific antibody and prevents the antibody binding to native streptokinase. The specification also describes a synthetic polypeptide (P1) comprising an epitope which binds to an streptokinase-specific antibody and reduces thrombolytic activity of streptokinase. P1 is used in thrombolytic therapy, and to prevent or treat glomerulonephritis and rheumatic fever.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antibody; epitope; trypsin; human; fibrinogen; detection; enzyme; cleavage; thrombosis; thrombus; thrombolysis.
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                                                                                                                            New polypeptides which bind streptokinase-specific antibodies useful in thrombolytic therapy
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Parhami-Seren B, Reed GL
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                                                                 WPI; 1999-190113/16.
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Matches 2; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Antinflammatory, T20/DP178; gp41 ectodomain; HIV-1 fusion; formyl peptide receptor family; FPR; inflammatory response up-regulation; chemoattractant.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present sequence is a peptide fragment of T20/DP178. T20/DP178 is a helical segment of the ectodomain of HIV-1 protein gp41. T21/DP178 is located in the carboxyl terminns of the gp41 ectodomain. gp41 plays a critical role in the fusion of HIV-1 and host cell membranes. T20/DP178 interacts with members of the formyl peptide receptor (FPR) family and thereby up-regulates an inflammatory response, and acts as a potent chemoattractant and activator of human peripheral blood phagocytes (but not T cells). The present peptide can be used to modulate an inflammatory response in a subject.
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   65.2%; Score 15; DB 20; Length 5; 100.0%; Pred. No. 7.8e+05; ive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                            AAB52657 standard; Peptide; 5 AA.
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4.405 Million cell updates/sec
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1: /cgn2_6/ptodata/1/pubpaa/USO8 NEW PUB.pep:*

2: /cgn2_6/ptodata/1/pubpaa/PCT_NEW FUB.pep:*

3: /cgn2_6/ptodata/1/pubpaa/USO6_NEW FUB.pep:*

4: /cgn2_6/ptodata/1/pubpaa/USO6_NEW PUB.pep:*

5: /cgn2_6/ptodata/1/pubpaa/USO6_NEW PUB.pep:*

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7: /cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*

8: /cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*

9: /cgn2_6/ptodata/1/pubpaa/USO8_PUBCOMB.pep:*

10: /cgn2_6/ptodata/1/pubpaa/USO9_NEW PUB.pep:*

11: /cgn2_6/ptodata/1/pubpaa/USO9_NEW PUB.pep:*

12: /cgn2_6/ptodata/1/pubpaa/USO9_NEW PUB.pep:*

13: /cgn2_6/ptodata/1/pubpaa/USO9_NEW PUB.pep:*

14: /cgn2_6/ptodata/1/pubpaa/USO9_NEW PUB.pep:*

15: /cgn2_6/ptodata/1/pubpaa/USO9_NEW PUB.pep:*
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6/ptodata/1/pubpaa/US60_PUBCOMB.pep:*
                  GenCore version 5.1.3
(c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-096-749A-94
US-09-866-135-1
US-08-444-409-35
US-08-424-550B-469
US-09-995-749A-14
US-09-764-884-2
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US-09-281-717-3

US-09-441-132-9

US-09-968-561A-211

US-09-968-561A-211

US-09-201-396-7

US-10-105-930-70

US-10-105-930-70

US-09-903-412-51

US-09-903-412-51
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                                                                                                                                                                                                                                                                                                                                                    140259 segs, 25548876 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries
                                                                                                                                     February 12, 2003, 10:52:34
                                                                                              protein search, using sw model
                                                                                                                                                                                                                                                                                          BLOSUM62
Gapop 10.0 , Gapext 0.5
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23
1 FXXLW 5
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                                        Copyright
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /cgn2_
                                                                                                                                                                                                                                                                                                                                                                                                                               seq length: 5 seq length: 5
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Match ]
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Perfect score:
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Maximum DB E
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GENERAL INFORMATION:

APPLICANT: Baxter, John D.

APPLICANT: Baxter, John D.

APPLICANT: Baxter, John D.

APPLICANT: Perg, Weijun

APPLICANT: Fletterick, Robert J.

APPLICANT: Fletterick, Robert J.

APPLICANT: Wagner, Richard L.

APPLICANT: Wagner,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ..
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Sequence 80, 1
Sequence 82, 2
Sequence 84, 3
Sequence 84, 3
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APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Picksley, Stephen
APPLICANT: Chene, Partick
APPLICANT: Chene, Partick
APPLICANT: Garcia-Excheverria, Carlos
APPLICANT: Garcia-Excheverria, Carlos
APPLICANT: Ruret, Pascal
TILLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REPERENCE: 4-2037/A/PCT
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT APPLICATION NUMBER: US/09/214,371B
FRIOR APPLICATION NUMBER: PCT/EP97/03549
FRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
82.6%; Score 19; DB 10; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 5; Conservative 0; Mismatches 0; Indels
US-09-903-412-69
US-09-903-412-74
US-09-903-412-76
US-09-903-412-78
US-09-903-412-82
US-09-903-412-82
US-09-903-412-84
US-09-903-412-84
                                                                                                                                                                                                                                                                                                                                                                 ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 2, Application US/09214371B; Patent No. US20010018511A1; GENERAL INFORMATION:
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Patent No. US20020061539A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
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US-09-214-371-2
       SEQ ID NO 2
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                                                                                                                                                                                                                                                                                                     .;
0
                                                                                                                                                                                                                                                 Query Match 82.6%; Score 19; DB 10; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 5; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              D377/AP6D
                                                                                                                                                           LOCATION: (2)..(3)
OTHER INFORMATION: Xaa = Any Amino Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5527.PC.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: JOHN N. SIMONS
APPLICANT: JOHN N. SIMONS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. CHLAUDER
APPLICANT: GEORGE G. CHLAUDER
APPLICANT: THOMAS P. LERRY
APPLICANT: ATHOMAS P. LERRY
APPLICANT: JAMES C. ERKER
APPLICANT: JAMES C. ERKER
APPLICANT: SHERI L. BUIJK
APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B. NO
TITLE OF INVENTION: REAGENTS AND MET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 228, Application US/08424550B
; Patent No. US20020119447A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: ABBOTT LABORATORIES
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: 435435
ATTORNEY/AGENT INFORMATION:
NAME: POREMESKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PF
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-938-2655
TELEFAX: 708-938-2655
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
                 SEQ ID NO 3
LENGTH: 5
TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
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                                                                                                                                  NAME/KEY: SITE LOCATION: (2)...
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US-08-424-550B-228
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APPLICANT: Tonlinson, Ian M
APPLICANT: Winter, Gregory
TITLE OF INVENTION: Method to Screen Phage Display Libraries with Different Ligands
FILE REFERENCE: 8039/1073B
CURRENT APPLICATION NUMBER: US/09/968,561A
CURRENT FILING DATE: 2001-10-01
PRIOR APPLICATION NUMBER: US 60/065,248
PRIOR FILING DATE: 1997-11-21
PRIOR FILING DATE: 1998-10-20
PRIOR FILING DATE: 1998-10-20
PRIOR FILING DATE: 1998-10-20
PRIOR FILING DATE: 1998-10-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Murphy, Andrew J.
APPLICANT: Paul, Joeremy
TITLE OF INVENTION: MFFREDS AND COMPOSITIONS FOR IDENTIFYING RECEPTOR
TITLE OF INVENTION: MFFRECTORS
FILE REPERENCE: CPI-012CP9
CURRENT FILING DATE: 1998-11-30
RARLIER APPLICATION NUMBER: 08/582,333
EARLIER APPLICATION NUMBER: 08/322,137
EARLIER PELING DATE: 1994-10-13
EARLIER PELING DATE: 1994-01-31
EARLIER PILING DATE: 1994-01-31
EARLIER PILING DATE: 1994-01-31
EARLIER APPLICATION NUMBER: 08/190,328
EARLIER PILING DATE: 1993-03-31
EARLIER PILING DATE: 1993-03-31
NUMBER OF SEQ ID NOS: 47
NUMBER OF SEQ ID NOS: 47
NUMBER OF SEQ ID NOS: 47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
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50.0%; Pred. No. 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1; Mismatches
                                        Sequence 217, Application US/09968561A; Patent No. US20020164642A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7, Application US/09201396A, Publication No. US20030009022A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQ ID NOS: 350
SOFTWARE: PatentIn version 3.1
SEQ ID NO 217
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US-09-968-561A-217
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                    US-09-968-561A-217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  4 LW 5
... 5
MW 5
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US-09-201-396-7
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LENGTH: 5
TYPE: PRT
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APPLICANT: Minter, Gregory
TILLE OF INVENTION! Method to Screen Phage Display Libraries with Different Ligands
FILE REFERENCE: 8039/1073B
CURRENT FILING DATE: 2001-10-01
FRIOR PAPLICATION NUMBER: US 60/968,561A
FRIOR APPLICATION NUMBER: US 60/065,248
FRIOR FILING DATE: 1997-11-21
FRIOR FILING DATE: 1997-11-21
FRIOR FILING DATE: 1997-11-21
FRIOR FILING DATE: 1997-11-21
FRIOR FILING DATE: 1997-11-31
FRIOR FILING DATE: 1997-11-31
FRIOR FILING DATE: 1997-11-31
FRIOR FILING DATE: 1997-11-31
FRIOR FILING DATE: 1998-10-20
FRIOR FILING DATE: 2000-02-24
NUMBER OF SEQ ID NOS: 350
FRIOR FILING DATE: 2000-02-24
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                                                                                                                                                           Sequence 1. Sequence 1. Sequence 1. Sequence 2. Patent No. US20020061848al |
GENERAL INCORMATION:
GENERAL INCORMATION:
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION FILE REFERENCE: 210121.469C8 |
CURRENT FILING DATE: 2001-04-23 |
NUMBER OF SEQ ID NOS: 599 |
SOFTWARE FESTSEQ for Windows Version 3.0/4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 211, Application US/09968561A Patent No. US20020164642A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-09-841-132-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
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; ORGANISM: Homo sapiens
US-09-968-561A-211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
Matches 1; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 2; Conserv
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                                                                                                                                            US-09-841-132-9
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56.5%; Score 13; DB 9; Length 5;

Query Match

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Gaps
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                                                                                                                                                                                                                                  US-10-105-930-70

| Sequence 70, Application US/10105930
| Sequence 70, Application US/10105930
| Publication No. US20030009018A1
| GENERAL INFORMATION: US20030009018A1iko
| APPLICANT: Maeda, Masatsugu
| APPLICANT: Maguchi, No. US20030009018A1iko
| TITLE OF INVENTION: NOVEL HENOPOIETIN RECEPTOR PROTEIN, NR12
| FILE REFERENCE: 06501-105US1
| CURRENT FILING DATE: 2002-03-25
| PRIOR APPLICATION NUMBER: PCT/JP00/06654
| PRIOR APPLICATION NUMBER: PCT/JP00/06654
| PRIOR APPLICATION NUMBER: JP 2000-240397
| PRIOR PILING DATE: 2000-08-03
| PRIOR FILING DATE: 1999-09-27
| NUMBER OF SEQ ID NOS: 77
| SOFTWARE FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             h Similarity 50.0%; Pred. No. 1.2e+05; 1; Conservative 1; Mismatches 0; Indels
ilarity 50.0%; Pred. No. 1.2e+05;
Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 56.5%; Score 13; DB 9; Length 5; Best Local Similarity 50.0%; Pred. No. 1.2e+05; Matches 1; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 51, Application US/09903412
FUDDIcation No. US20030027319A1
GENERAL INFORMATION:
APPLICANT: Koide, Shohei
TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
FILE REPERENCE: 109.050US1
CURRENT APPLICATION NUMBER: US/09/903,412
FRIOR APPLICATION NUMBER: US 60/217,474
FRIOR FILING DATE: 2000-07-11
FRIOR APPLICATION NUMBER: US 60/217,474
NUMBER OF SEQ ID NOS: 121
SOPTWARE: FRASESO for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-105-930-70
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SEQ ID NO 51
                                                                                                            3
WM :-
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2 MW 3
                                                                                4 LW 5
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Gaps
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APPLICANT: Winter, Greg
APPLICANT: Tomlinson, Ian
TILLE OF INVENTION: Methods for Selecting Functional Peptides
FILE REFERENCE: 3789/72916
CURRENT APPLICATION NUMBER: 08/09/192,854
CURRENT FILING DATE: 1998-11.17
EARLIER FILING DATE: 1997-11.21
NUMBER OF SEQ ID NOS: 212
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 56.5%; Score 13; DB 10; Length 5; Best Local Similarity 50.0%; Pred. No. 1.2e+05; Matches 1; Conservative 1; Mismatches 0; Indels
                                                                                                  APPLICANT: Koieda, Shohei
TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SOUDRNCES: 118
CORRESPONDENCE ADDRESS:
ADDRESSER: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Bighth Street, Ste. 1600
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                109.034US1
                                                                                                                                                                                                                                                                                                                                                                                                                                             US/09/096,749A
                                     Sequence 51, Application US/09096749A
Patent No. US20020019517A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 120, Application US/09192854
; Patent No. US20020068276A1
                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATORNEY AGENT:
NAME: Ann S. Viksnins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.0
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
TELEPAX: (612) 373-6900
TELEPAX: (612) 373-6900
SEQUENCE CHARACTERISTICS: 51:
                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/09/C
FILING DATE: June 12, 1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM Compatible OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                        Diskette
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                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
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STRANDEDNESS: sin
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                                                                                                                                                                                                                                                                         USA
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RESULT 10
US-09-096-749A-51
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                                                                                                                                                                                                                                                                         COUNTRY:
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## APPLICANT: TRINCHIER; GIORGIO
## APPLICANT: TRINCHIER; GIORGIO
## APPLICANT: TRINCHIER; GIORGIO
## APPLICANT: CARK, STEVEN C.
## APPLICANT: GORDON G.
## APPLICANT: MONG, GORDON G.
## APPLICANT: MONGER: US/09/924, 703
## TITLE OF INVENTION: ANTIBODIES TO NATURAL KILLER STIMULATORY FACTOR
## FILE REFERENCE: 01142.01100
## CURRENT FILING DATE: 1999-06-04
## PRIOR APPLICATION NUMBER: 08/828,000
## PRIOR FILING DATE: 1997-05-16
## PRIOR FILING DATE: 1995-03-13
## PRIOR FILING DATE: 1996-09-18
## PRIOR PELING DATE: 1989-0-09-18
## PRIOR APPLICATION NUMBER: 07/307,817
## PRIOR APPLICATION NUMBER: 07/307,817
## PRIOR APPLICATION NUMBER: 07/307,817
## PRIOR PELING DATE: 1988-01-10
## PRIOR FILING DATE: 1988-11-10
## NUMBER OF SEQ ID NOS: 19
## COFTWARE PARCELIN VET: 2.1
                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                               ;
0
                                                                      Query Match 56.5%; Score 13; DB 10; Length 5; Best Local Similarity 50.0%; Pred. No. 1.2e+05; Matches 1; Conservative 1; Mismatches 0; Indels
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STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Savakis, Charalambos
APPLICANT: Savakis, Charalambos
APPLICANT: Franz, Gerald H.
APPLICANT: Loukeris, Athanasios
TITLE OF INVENTION: Eukaryotic Transposable Element
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 4, Application US/08239765B
Patent No. US20020026650A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                 Sequence 9, Application US/09924703
Patent No. US20020137898A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
; ORGANÎSM: Homo sapiens
US-09-192-854-120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: Two Mili
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-239-765B-4
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US-09-06-749A-94
Sequence 94, Application US/09096749A
Sequence 94, Application US/09096749A
Setent No. US20020019517A1
GENERAL INFORMATION:
TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, Ste. 1600
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 5;
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                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/239,765B
FILING DATE: 09-MAY-1994
CLASSIFICATION 1435
PRIOR APPLICATION 1435
PRIOR APPLICATION NUMBER: US 07/946,237
FILING DATE: 14-SEP-1992
ATTONNET/AGENT INFORMATION:
NAME: BATCOK, David B:
REGISTRATION NUMBER: 22,592
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 12; DB 8; I
Pred. No. 1.2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Mismatches
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COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/096,749A
FILING DATE: June 12, 1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NIMBER: 109.034US1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
TELEFAX: (612) 339-3061
                                                                                                                                                                                        COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Ann S. Viksnins
REGISTRATION NUMBER: 37,748
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 52.2%;
Best Local Similarity 50.0%;
Matches 1; Conservative
                                                                                                                                                       Floppy disk
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            , MOLECULE TYPE: peptide US-08-239-7658-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                              COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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                                                                         02173
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COUNTRY:
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compacible
OPERATIES: SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: ABBOTT LABORATORIES D377/AP6D 100 ABBOTT PARK ROAD
                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 690068.409C1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (206) 622-4900
TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/484,409
FILLING DATE: 07-UNN-1995
CLASSIFTCATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: JOHN N. SIMONS
APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: STRESH M. DESAI
APPLICANT: THOMAS P. LERRY
APPLICANT: ANTHONY SCOTT MUERHOFF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 469, Application US/08424550B patent No. US20020119447A1 GENERAL INFORMATION:
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ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: Maki, David J.
REGISTRATION NUMBER: 31,392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JAMES C. ERKER
SHERI L. BUIJK
ISA K. MUSHAHWAR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 amino acids
                                                                                  COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                       Washington
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nes 1; Conserv
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                                                               98104-7092
Seattle
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APPLICANT:
APPLICANT:
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APPLICANT: PTITSTN, Leonid R
APPLICANT: PTITSTN, Leonid R
APPLICANT: PTITSTN, Leonid R
APPLICANT: SMIRNOV, Sergey V
APPLICANT: SMIRNOV, Yulia G
APPLICANT: ROSTOVA, Tatyana A
APPLICANT: TAMPOLSKAYA, Tatyana V
APPLICANT: MARDAINER, Mikhail M
TITLE OF INVENTION: NEW MUTANT-ACETYLGLUTAMATE SYNTHASE AND METHOD FOR L-ARGININE PRC
FILE REFRERENCE: 209873UGO
CURRENT FILING DATE: 2001-06-22
PRIOR APPLICATION NUMBER: RUSSIAN FEDERATION 2000116481
PRIOR PLING DATE: 2000-06-28
PRIOR PLING DATE: 2000-06-28
PRIOR PLING DATE: 2000-06-28
PRIOR PLING DATE: 2000-06-28
PRIOR PLING DATE: 2000-06-36
NUMBER OF SEC ID NOS: 14
SOFTMARE: PATENTIN VERSION 3.1
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Patent No. US20020076412A1

GENERAL INFORMATION:

APPLICANT: Steinman, Lawrence
APPLICANT: Zamvil, Scott

TITLE OF INVENTION: METHODS FOR MODULATING THE IMMUNE SYSTEM
NUMBER OF SEQUENCES: 52

CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         52.2%; Score 12; DB 10; Length 5; 50.0%; Pred. No. 1.2e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                      52.2%; Score 12; DB 10; Length 5; 50.0%; Pred. No. 1.2e+05; tive 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 1, Application US/09866135
Patent No. US20020051443A1
  94:
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les 1; Conservative
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Best Local Similarity 50.09
Matches 1; Conservative
                     SEQUENCE CHARACTERISTICS:
  INFORMATION FOR SEQ ID NO:
                                        LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                 internal
                                                                                            Sirker TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                          ANTI-SENSE: NO
FRAGMENT TYPE: i
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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LENGTH: 5
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47.8%; Score 11; DB 9; Length 5; 100.0%; Pred. No. 1.2e+05; tive 0; Mismatches 0; Indels
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; Patent No. US20020160949A1
                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 15, Application US/09895593 ; Patent No. US20020160949A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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APPLICANT: Pandey, Akhilesh
APPLICANT: Ozaki, Katsutoshi
                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Pandey, Akhilesh
APPLICANT: Ozaki, Katsutoshi
                                                                                                                                                                Best Local Similarity 100. Matches 1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Baumann, Heinz
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                                  NAME/KEY: Site
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APPLICANT:
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APPLICANT:
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APPLICANT:
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Fatent No. US20020161208A1;
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
CURRENT APPLICATION NUCLEIC Acids, Proteins, and Antibodies
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper;
NUMBER OF SEQ ID NOS: 38
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                               ;
                                                                                                                                                                                                                                                                                              47.8%; Score 11; DB 8; Length 5; 100.0%; Pred. No. 1.2e+05; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 11, Application US/0995749A
Patent No. US20020155568A1
GENERAL INFORMATION:
APPLICANT: VAN GEEL-SCHUTTEN, GERRITDINA HENDRIKA
APPLICANT: DIJKHUIZEN, LUBBERT
APPLICANT: LEER, ROBERT-JAN
TITLE OF INVENTION: NOVEL GLUCOSYLTRANSFERASES
FILE REFERENCE: B043388-CIP
CURRENT APPLICATION NUMBER: US/09/995,749A
CURRENT FILING DATE: 2001-11-29
FRIOR FILING DATE: 2000-6-28
FRIOR FILING DATE: 2000-06-28
FRIOR FILING DATE: 2000-05-25
                              5527.PC.01
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; ORGANISM: Neisseria polysaccharea
US-09-995-749A-14
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 55:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-938-2623
TELEFAK: 708-938-2623
                                                                                      TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 469:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                            1; Conservative
                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: protein
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ORGANISM: Homo sapiens
                                                                                                                                                                                amino acid
XGY: linear
                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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Best Local Similarity
Matches 1; Conserv
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US-09-995-749A-14
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LENGTH: 5
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53
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. LOCATION: (3)
. OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
US-09-764-884-2
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APPLICANT: Farr, Andrew G.
APPLICANT: Erin, Steven F.
APPLICANT: Leonard, Warren G.
APPLICANT: Leonard, Warren J.
APPLICANT: Leonard, Warren J.
APPLICANT: Leonard, Warren J.
TITLE OF INVENTION: Thymic Stromal Lymphopoietin Receptor Molecules and TITLE OF INVENTION: Uses Thereof
FILE REFERENCE: 00-514-E
CURRENT APPLICATION NUMBER: US/09/895,593
CURRENT APPLICATION NUMBER: 60/215,658
PRIOR PILING DATE: 2000-06-28
NUMBER OF SEQ ID NOS: 16
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 15
LENGTH: 5
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APPLICANT: Levin, Steven D.
APPLICANT: Earr, Andrew G.
APPLICANT: Ziegler, Steven F.
APPLICANT: Leonard, Warren J.
APPLICANT: Leonard, Warren J.
APPLICANT: Looish, Harvey F.
TITLE OF INVENTION: Thymic Stromal Lymphopoietin Receptor Molecules and TITLE OF INVENTION: Uses Thereof
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APPLICANT: Rosen et al.
TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REPRENCE: P7211
CURRENT APPLICATION NUMBER: US/09/764,857
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper NUMBER OF SEQ ID NOS: 13
                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: SATO, KOH
APPLICANT: TSUMENARI, TOSHIAKI
APPLICANT: TSUMENARI, TOSHIAKI
TITLE OF INVENTION: CACHEXIA REMEDY
FILE REPERBROKE: 04833-0038
CURRENT APPLICATION NUMBER: US/09/423,800
CURRENT FILING DATE: 1999-11-12
PRIOR APPLICATION NUMBER: PCT/JP98/02116
PRIOR PLILING DATE: 1998-05-13
PRIOR PLILING DATE: 1997-05-15
PRIOR PLILING DATE: 1997-05-15
PRIOR PLILING DATE: 1997-07-18
             CURRENT APPLICATION NUMBER: US/09/895,593
CURRENT FILING DATE: 2001-06-28
PRIOR APPLICATION NUMBER: 60/215,658
PRIOR FILING DATE: 2000-06-28
NUMBER OF SEQ ID NOS: 16
SOFTWARE: PatentIn Ver. 2.0
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; Patent No. US20020165363A1
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Patent No. US20020164685A1
                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
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FILE REFERENCE: 00-514-E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Homo sapiens
US-09-423-800-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: SATO, K
                                                                                                                                                                                                                                                                                                                                                                         US-09-895-593-16
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US-09-764-857-2
                                                                                                                                                                       SEQ ID NO 16
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; OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids US-09-860-670-2
                                                                                                                                                                                    OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
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; Sequence 2, Application US/09860670
; Patent No. US20020165137A1
; GENERAL INFORMATION:
; APPLICANT: Ruben et al.
; TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
; FILE REFERENCE: PA12TION NUMBER: US/09/860,670
; CURRENT APPLICATION NUMBER: US/09/860,670
; CURRENT APPLICATION NUMBER: US/09/860,670
; CURRENT APPLICATION NUMBER: 201-05-21
; Prior application data removed - consult PALM or file wrapper
; NUMBER OF SEQ ID NOS: 289
; SOFTWARES PATENTIN Ver. 2.0
; SEQ ID NO 2
: LIENGTH: 5
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TITLE OF INVENTION: 70 Human Secreted Proteins
FILE REFERENCE: PZ001P1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT FILING DATE: 2001-10-19
CURRENT FILING DATE: 2001-10-19
PRIOR APPLICATION NUMBER: 09/148,545
PRIOR PELING DATE: 1998-09-04
PRIOR PELING DATE: 1998-09-04
PRIOR PELING DATE: 1997-03-07
PRIOR APPLICATION NUMBER: 60/040,162
PRIOR FILING DATE: 1997-03-07
PRIOR PELING DATE: 1997-03-07
PRIOR PELING DATE: 1997-03-07
PRIOR PELING DATE: 1997-03-07
PRIOR APPLICATION NUMBER: 60/040,161
PRIOR APPLICATION NUMBER: 60/040,626
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; Patent No. US20020164669A1
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2
LENGTH: 5
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ORGANISM: Homo sapiens
                                                                   TYPE: PRT
ORGANISM: Homo sapiens
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nes 1; Conserv
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NAME/KEY: Site
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LOCATION: (3)
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US-09-981-876-2
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FILING DATE: 1997-03-07
APPLICATION NUMBER: 60/040,163
FILING DATE: 1997-03-07
APPLICATION NUMBER: 60/047,615
FILING DATE: 1997-05-23 FILING DATE: 1997-03-07 APPLICATION NUMBER: 60/040,334 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,587 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,582 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,596 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,632 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,601 APPLICATION NUMBER: 60/043,312 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,313 FILING DATE: 1997-04-11 FILING DATE: 1997-03-07 APPLICATION NUMBER: 60/040,336 APPLICATION NUMBER: 60/047,600 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,597 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,633 APPLICATION NUMBER: 60/047,583 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,617 APPLICATION NUMBER: 60/047,618 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,503 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,592 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,584 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,492 APPLICATION NUMBER: 60/047,598 APPLICATION NUMBER: 60/047,612 APPLICATION NUMBER: 60/043,580 APPLICATION NUMBER: 60/043,568 APPLICATION NUMBER: 60/043,314 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,569 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,311 APPLICATION NUMBER: 60/043,674 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/043,672 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/047,502 APPLICATION NUMBER: 60/047,581 APPLICATION NUMBER: 60/047,500 APPLICATION NUMBER: 60/047,613 APPLICATION NUMBER: 60/043,671 FILING DATE: 1997-04-11 71LING DATE: 1997-05-23 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 LING DATE: 1997-04-11 FILING DATE: 1997-05-23 ILING DATE: 1997-05-23 ILING DATE: 1997-04-11 FILING DATE: 1997-04-1 FILING DATE: 1997-04-1 ILING DATE:

R FILING DATE: 1997-08-22

R APPLICATION NUMBER: 60/056,892

R FILING DATE: 1997-08-22

R APPLICATION NUMBER: 60/047,595

R FILING DATE: 1997-05-23

R APPLICATION NUMBER: 60/057,761 FILING DATE: 1997-04-11 APPLICATION NUMBER: 60/048,974 FILING DATE: 1997-06-06 APPLICATION NUMBER: 60/056,886 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,877 FILING DATE: 1997-08-22
APPLICATION UNMBER: 60/056,889
FILING DATE: 1997-08-22
APPLICATION NUMBER: 60/056,893 NUMBER: 60/043,315 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,878 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,911 FILING DATE: 1997-08-22 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,614 APPLICATION NUMBER: 60/056,630 APPLICATION NUMBER: 60/056,662 FILING DATE: 1997-08-22 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,882 APPLICATION NUMBER: 60/056,888 FILING DATE: 1997-08-22 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,874 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,910 FILING DATE: 05-Sep-1997 APPLICATION NUMBER: 60/047,599 APPLICATION NUMBER: 60/056,872 APPLICATION NUMBER: 60/056,637 APPLICATION NUMBER: 60/056,903 APPLICATION NUMBER: 60/056,879 APPLICATION NUMBER: 60/056,880 APPLICATION NUMBER: 60/056,894 APPLICATION NUMBER: 60/056,636 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,864 APPLICATION NUMBER: 60/056,631 FILING DATE: 1997-08-22 APPLICATION NUMBER: 60/056,845 FILLING DATE: 1997-05-23 APPLICATION NUMBER: 60/047,588 APPLICATION NUMBER: 60/047,585 APPLICATION NUMBER: 60/047,586 APPLICATION NUMBER: 60/047,590 APPLICATION NUMBER: 60/047,594 APPLICATION NUMBER: 60/047,589 APPLICATION NUMBER: 60/047,593 FILING DATE: 1997-05-23 APPLICATION NUMBER: 60/043,578 FILING DATE: 1997-08-22 FILING DATE: 1997-08-22 1997-08-22 1997-08-22 FILING DATE: 1997-05-23 1997-05-23 1997-05-23 FILING DATE: 1997-05-23 FILING DATE: 1997-05-23 1997-05-23 FILING DATE: FILING DATE: FILING DATE: APPLICATION PRIOR PRIOR

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| PRIOR FILING DATE: 1997-04-1, 50
| PRIOR FILING DATE: 1997-04-1, 50
| PRIOR PILING DATE: 1997-06-20-6, 64
| PRIOR PILING DATE: 1997-06-20-6, 67
| PRIOR PILING DATE: 1997-06-20-6, 66
| PRIOR PILING DATE: 1997-06-20-6, 67
| PRIOR PILING DATE: 1997-06-20-6, 684
| PRIOR PILING DATE: 1997-06-20-6, 67
| PRIOR PILING DATE: 1997-06-20-6, 684
| PRIOR PILING DATE: 1997-06-20-6, 694
| PRIOR PILING DATE: 1997-06-20-6, 694
| PRIOR PILING DATE: 1997-06-20-6, 694
| PRIOR PILING DATE: 1997-06-20-19-6, 694
| PRIOR PILING DATE: 1997
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OTHER INFORMATION: Xaa equals any of the L-amino acids commonly found in naturally OTHER INFORMATION: proteins
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TITLE OF INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PT270
CURRENT APPLICATION NUMBER: US/09/764,868
CURRENT FILING DATE: 2001-01-17
Prior application data removed - refer to PALM or file wrapper NUMBER OF SEQ ID NOS: 1510
SOFTWARE: PatentIn Ver. 2.0
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PRIOR APPLICATION NUMBER: US 60/048,160
PRIOR PRILING DATE: 1997-05-30
PRIOR PRILING DATE: 1997-05-30
PRIOR PRILING DATE: 1997-05-30
PRIOR FILING DATE: 1997-05-30
PRIOR FILING DATE: 1997-05-30
PRIOR FILING DATE: 1997-05-30
PRIOR APPLICATION NUMBER: US 60/054,804
PRIOR FILING DATE: 1997-08-19
PRIOR PRILING DATE: 1997-08-19
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PRIOR FILING DATE: 1997-10-02
NUMBER OF SEQ ID NOS: 343
SEQ ID NO 2
LENGTH: 5
LENGTH: 5
                        PILING DATE: 1997-05-30
APPLICATION NUMBER: US 60/050,937
FILING DATE: 1997-05-30
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APPLICATION NUMBER: US 60/048,352
FILING DATE: 1997-05-30
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APPLICATION NUMBER: US 60/048,069
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FILING DATE: 1997-05-30
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FILING DATE: 1997-05-30
APPLICATION NUMBER: US 60/048,135
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ORGANISM: Homo sapiens
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LENGTH: 5
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NAME/KEY: Site
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US-09-764-904-2
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Patent No. US20020168684A1

GENERAL INFORMATION:

APPLICANT: CCMB, Michael J.

APPLICANT: CCMB, Michael J.

APPLICANT: CMB, Michael J.

APPLICANT: CMB, Michael J.

APPLICANT: CMB, Michael J.

APPLICANT: CMB, Michael J.

TITLE OF INVENTION: PRODUCTION OF MOTIF-SPECIFIC AND CONTEXT-INDEPENDENT ANTIBODIES U.

TITLE OF INVENTION: PREPRIED LIBRARIES AS ANTIGENS

TITLE OF INVENTION: PREPRIED LIBRARIES AS ANTIGENS

TITLE OF INVENTION: DATE: 2002-03-18

CURRENT APPLICATION NUMBER: US 09/148,712

PRIOR APPLICATION NUMBER: US 09/148,712

PRIOR PILING DATE: 1998-09-04

PRIOR FILING DATE: 2000-03-24

NUMBER OF SEQ ID NOS: 145

SOFFWARE PATENTIN OF SECOND 3.1
; NAME/KEY: Site
; CCATION: (3)
; CTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
US-09-764-868-2
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LOCATION: (3)..(3)

1 THER INCRMATION: PHOSPHORYLATION; threonine at position 3 is phosphorylated US-10-014-485A-79
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Sequence 42, Application US/10104019
Sequence 42, Application US/10104019
Sequence 42, Application
Sequence 4
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                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                      Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 28
US-10-014-485A-79
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US-10-104-019-42
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. LOCATION: (3)
; OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
US-09-764-904-2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 2, Application US/09764904;
Sequence 2, Application US/09764904;
Patent No. US20020173454A1
GENERAL INFORMATION:
APPLICANT: Rosen et al.
TITLE OP INVENTION: Nucleic Acids, Proteins, and Antibodies
FILE REFERENCE: PA122
CURRENT FILING DATE: 2001-01-17
Prior application data removed - consult PALM or file wrapper
NUMBER OF SEQ ID NOS: 137
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 2
LENGTH: 5
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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Sequence 2, Application US/09852797
Parent No. US20020172994A1
GENERAL INFORMATION:
APPLICANT: ROSEN et al.
TILE OF INVENTION: 28 Human Secreted Proteins
FILE REPERENCE: P2003P2
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CURRENT FILING DATE: 2001-05-11
PRIOR APPLICATION NUMBER: 60/265,583
PRIOR FILING DATE: 2001-02-02
PRIOR PILING DATE: 1998-09-11
PRIOR FILING DATE: 1998-03-11
PRIOR FILING DATE: 1998-03-12
PRIOR FILING DATE: 1998-03-12
PRIOR FILING DATE: 1998-03-12
PRIOR PLING DATE: 1998-03-14
PRIOR PLING DATE: 1997-03-14
PRIOR FILING DATE: 1993-10-05
PRIOR APPLICATION WUMBER: 07/969,071
PRIOR FILING DATE: 1992-10-30
NUMBER OF SEQ ID NOS: 108
SOFTWARE: PATENTIN Ver. 2.1
SEQ ID NO 42
LENGTH: 5
                                                                                                                                                                                               TYPE: PRT ORGANISM: Artificial Sequence
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ORGANISM: Homo sapiens
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Sequence 8, Application US/10035349
; Sequence 8, Application US/10035349
; Patent No. US20020172961A1
; GENERAL INFORMATION:
; APPLICANT: Schneider, Luke V.
; APPLICANT: Hall, Michael P.
; APPLICANT: Target Discovery, Inc.
; APPLICANT: Target Discovery, Inc.
; TITLE OF INVENTION: Sequences
; TITLE OF INVENTION: Sequences
; TITLE OF INVENTION: Sequences
; TITLE OF INVENTION: US Sequences
; FILE REFERENCE: 02044-00071008
; CURRENT FILING DATE: 2002-04-15
; PRIOR APPLICATION NUMBER: US 60/242,165
; PRIOR APPLICATION NUMBER: US 60/242,398
; PRIOR FILING DATE: 2000-10-19
; NUMBER OF SEQ ID NOS: 13
; SEQ ID NO 8
; LENGTH: 5
                                                                                                                                                                                          APPLICANT: Petersch, Robert
APPLICANT: Target Discovery, Inc.
APPLICANT: Target Discovery, Inc.
TITLE OF INVENTION: Mass Defect Labeling for the Determination of Oligomer TITLE OF INVENTION: Sequences
FILE REPRESENCE: 0.2044-0.00710US
CURRENT APPLICATION NUMBER: US 60/242,165
PRIOR APPLICATION NUMBER: US 60/242,165
PRIOR APPLICATION NUMBER: US 60/242,398
PRIOR FILING DATE: 2000-10-19
PRIOR FILING DATE: 2000-10-19
RIOR FILING DATE: 2000-10-19
RIOR FILING DATE: 2000-10-19
RIOR FILING DATE: 2000-10-19
ROWBER OF SEQ ID NOS: 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Xaa = Gly labeled with 5-bromo-3-pyridylacetic OTHER INFORMATION: acid (5-Br-3-PAA)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                   ; Sequence 2, Application US/10035349; Patent No. US20020172961A1
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ORGANISM: Artificial Sequence
                                                                                                                                       APPLICANT: Schneider, Luke V. APPLICANT: Hall, Michael P.
                                                                                                               GENERAL INFORMATION:
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NAME/KEY: MOD RES
                                 US-10-035-349-2
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LENGTH: 5
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APPLICANT: Campos, Reinaldo
APPLICANT: Campos, Reinaldo
APPLICANT: No. US20020113633A10gaki, Hiroyuki
APPLICANT: No. US20020113633A10gaki, Hiroyuki
APPLICANT: Suslow, Trevor
APPLICANT: The Regents of the University of California
APPLICANT: The Regents of the University of Phenylalanine Ammonia-Lyase (PAL)
TITLE OF INVENTION: Gene in Wounded Lettuce Tissue
FILE REFERENCE: 023070-12450004
FILE REPERENCE: 023070-12450004
CURRENT APPLICATION NUMBER: US/09/964,992A
CURRENT APPLICATION NUMBER: US 60/235,956
PRIOR PLING DATE: 2000-09-26
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                                             PRIOR FILING DATE: 1997-05-30
PRIOR PILING DATE: 1997-05-30
PRIOR PILING DATE: 1997-05-30
PRIOR FILING DATE: 1997-05-30
PRIOR FILING DATE: 1997-05-30
PRIOR FILING DATE: 1997-05-30
PRIOR PILING DATE: 1997-05-30
PRIOR PILING DATE: 1997-05-30
PRIOR PLING DATE: 1997-09-65
PRIOR PELING DATE: 1997-09-65
PRIOR APPLICATION NUMBER: 60/048,970
PRIOR PELING DATE: 1997-09-06
PRIOR PILING DATE: 1997-09-06
PRIOR FILING DATE: 1997-12-19
NUMBER OF SEQ ID NOS: 118
SEQ ID NO 2
LENGTH: 5
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                                 APPLICATION NUMBER: 60/050,934
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FILING DATE: 1997-03-14
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SOFTWARE: PatentIn Ver. 2.1
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Matches 1; Conserva
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TITLE OF INVENTION: Apoptotic Protease Mch6, Nucleic Acids
Encoding Same and Methods of Use
                                                                                                                                                                                                    APPLICANT: Dinemri, Emad S. Perlandes-Alnemri, Teresa Litwack, Gerald Grandes Mché, Nucleic Acids TITLE OF INVENTION: Encoding Same and Methods of Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTY: United States
ZIE: 92122
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
COMPUTER: 1BM PC COMPA: DS SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/059,749
FILING DATE: 29-Jan-2002
CLASSIFICATION: Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/865,579
FILING DATE: 29-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                    CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Cambell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-ID 2180
TELECOMMUNICATION INFORMATION:
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell & Flores LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 63, Application US/10059749; Publication No. US20020183504A1; GENERAL INFORMATION: APPLICANT: Alnemri, Emad S.; Fernandes-Alnemri, Teresa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 57:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-9849
                                                                                                                              sequence 57, Application US/10059749; Publication No. US20020183504A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 5 amino acids
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                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 87
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
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                                                                                              RESULT 37
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US-09-828-272A-6
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Publication No. US20020183255A1
GENERAL INFORMATION:
APPLICANT: LIPTON, James M
APPLICANT: CATANIA, Anna P
ITILE OF INVENTION: USE OF KPV TRIPEPTIDE FOR DERMATOLOGICAL DISORDERS
TITLE OF INVENTION: USE OF KPV TRIPEPTIDE FOR DERMATOLOGICAL DISORDERS
TITLE OF INVENTION: USE OF KPV TRIPEPTIDE FOR DERMATOLOGICAL DISORDERS
CURRENT APPLICATION NUMBER: US/09/828,272A
CURRENT FILING DATE: 2001-04-06
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin version 3.1
SEQ ID NO 6
LENGTH: 5
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                                            0; Indels
47.8%; Score 11; DB 9; Length 5; 100.0%; Pred. No. 1.2e+05; tive 0; Mismatches 0; Indels
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US-10-042-141-2
; Sequence 2, Application US/10042141
; Publication No. US20020183503A1
; GENERAL INFORMATION:
; APPLICANT: Ruben et al.; TITLE OF INVENTION: 26 Human secreted proteins
; FILE REFERENCE: PZ040P1
; CURRENT APPLICATION NUMBER: US/10/042,141
; CURRENT APPLICATION NUMBER: O9/726,643
PRIOR APPLICATION NUMBER: O9/726,643
; PRIOR FILING DATE: 2000-01-01
; PRIOR PILING DATE: 2000-06-02
; PRIOR APPLICATION NUMBER: 60/137,725
; PRIOR APPLICATION NUMBER: 60/137,725
; NUMBER OF SEQ ID NOS: 190
; SEQ ID NOS: 190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ORGANISM: Artificial Sequence
                                                 1; Conservative
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  Query Match
Best Local Similarity
                                                                                                                                                                                                                RESULT 35
US-09-828-272A-6
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Publication No. US20020183504A1
GENERAL INFORMATION:
APPLICANT: Alnemri, Emad S.
Litwack, Gerald
TITLE OF INVENTION: Apoptotic Protease Mch6, Nucleic Acids
Encoding Same and Methods of Use
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/059,749
FILING DATE: 29-Jan-2002
CLASSIFICATION DATA:
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSE: Campbell & Flores LLP
STREET: 4770 La Jolla Village Drive, Suite 700
CITY: San Diego
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY AGENTE: 29-MAY-1997
ATTORNEY AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
FERERENCE/COCKET NUMBER: P-ID 2180
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/865,579
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FILING DATE: 29-MAY-1997
ATTORNEY/AGENT INFORMATION:
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APPLICATION NUMBER: US/10/059,749
FILING DATE: 29-Jan-2002
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 63:
US-10-059-749-63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (619) 535-9849
                                                                   COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 5 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 63:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS
                                                                                       ZIP: 92122
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 92122
COMPUTER READABLE FORM:
                                               STATE: California
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                           San Diego
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APPLICANT: Tartalia, Louis A.
APPLICANT: Tepper, Robert I.
APPLICANT: Culpeper, Janice A.
APPLICANT: Culpeper, Janice A.
APPLICANT: White, David W.
TITLE OF INVENTION: THE OB RECEPTOR AND METHODS FOR
TITLE OF INVENTION: THE DIAGNOSIS AND TREATMENT OF BODY WEIGHT DISORDERS,
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0
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47.8%; Score 11; DB 9; Le
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
COMPUTER: RANGOWS S
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENY APPLICATION DATA:
FILING DATE: 2002-FEB-19
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-ID 2180
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEPAX: (619) 535-9849
INFORMATION FOR SEQ ID NO: 69:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                             TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 69:
US-10-059-749-69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: 2002-FEB-19
FRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/864,564
FILING DATE: 28-MX-1997
APPLICATION NUMBER: 08/708,123
FILING DATE: 03-SEP-1996
APPLICATION NUMBER: 08/599,455
FILING DATE: 22-APR-1996
APPLICATION NUMBER: 08/599,455
FILING DATE: 22-DAN-1996
APPLICATION NUMBER: 08/591,455
FILING DATE: 28-DEC-1995
APPLICATION NUMBER: 08/570,142
FILING DATE: 11-DEC-1995
APPLICATION NUMBER: 08/59,455
FILING DATE: 11-DEC-1995
APPLICATION NUMBER: 08/570,142
FILING DATE: 11-DEC-1995
APPLICATION NUMBER: 08/569,485
FILING DATE: 08-DEC-1995
APPLICATION NUMBER: 08/569,485
FILING DATE: 08-DEC-1995
APPLICATION NUMBER: 08/569,485
FILING DATE: 08-DEC-1995
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APPLICATION WINBER: 08/562,663
FILING DATE: 27-NOV-1995
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 6, Application US/10079625; Publication No. US20020182676A1; GENERAL INFORMATION:
                                                                                                                                                                                                                         LENGTH: 5 amino acids
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US-10-079-625-6
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NAME: Meiklejohn, Ph.D., Anita L.

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APPLICANT: Browerda, Christian
APPLICANT: Browerda, Christian
APPLICANT: Browerda, Christian
APPLICANT: Browerda, Christian
TITLE OF INVENTION: Component of Stem Bromelain (as amended)
FILE REPERBENCE: 0623.0766002
CURRENT APPLICATION NUMBER: US/10/029,301
CURRENT FILING DATE: 1999-08-25
FRIOR APPLICATION NUMBER: DGT/GB98/00592
FRIOR APPLICATION NUMBER: CB 9704252.7
FRIOR APPLICATION NUMBER: GB 9704252.7
FRIOR PILING DATE: 1997-02-28
FRIOR FILING DATE: 1997-02-25
FRIOR FILING DATE: 1997-02-25
FRIOR FILING DATE: 1997-02-25
FRIOR FILING DATE: 1997-02-25
                                                                                                                                                                                                                                                                                                                                                             Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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TITLE OF INVENTION: METHODS OF TREATING BLADDER DISORDERS
FILE REFERENCE: 08191-022001
CURRENT APPLICATION NUMBER: US/10/074,956
CURRENT FILING DATE: 2002-06-10
PRIOR APPLICATION NUMBER: 60/268,175
                          07334/019002
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Publication No. US20020193332A1
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 4, Application US/10029301
Publication No. US20020188107A1
GENERAL INFORMATION:
                  REFERENCE/DOCKET NUMBER: 0733.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-542-6970
TELERAX: 617-542-8966
TELEX: 200154
INFORMATION FOR SEQ ID NO: 6: 52QUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
REGISTRATION NUMBER: 35,283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                       ; TOPOLOGY: unknown
; MOLECULE TYPE: peptide
US-10-079-625-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Ananas comosus
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TILLE OF INVENTION: Cytokine Receptor Common Gamma Chain Like
FILE REFERENCE: PF466P2
CURRENT APPLICATION WUMBER: US/10/078,059
CURRENT FILING DATE: 2002-02-20
FRIOR APPLICATION WUMBER: 60/269,876
PRIOR PLING DATE: 2000-03-21
PRIOR PLING DATE: 2000-08-17
PRIOR APPLICATION WUMBER: PCT/US00/22493
PRIOR PLING DATE: 1099-08-18
PRIOR PLING DATE: 1999-08-18
PRIOR PLING DATE: 1999-03-05
PRIOR PLING DATE: 1999-03-05
PRIOR PLING DATE: 1999-03-05
PRIOR PLING DATE: 1998-03-05
PRIOR PLING DATE: 1998-03-05
PRIOR PLING DATE: 1998-03-05
PRIOR PLING DATE: 1998-03-05
PRIOR APPLICATION WUMBER: 60/086,505
PRIOR PLING DATE: 1998-03-19
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APPLICANT: Paesen, Guido, Christiaan
TITLE OF INVENTION: Treatment of Allergic Rhinitis
FILE REFERENCE: 2488-1-004
CURRENT APPLICATION NUMBER: US/10/087,195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DITHER INFORMATION: Xaa equals any amino acid
                     NUMBER OF SEQ ID NOS: 29
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 6
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 5, Application US/10078059; Publication No. US20020193305A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 3, Application US/10087195; Publication No. US20020193306A1; GENERAL INFORMATION:
PRIOR FILING DATE: 2001-02-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Ver. 2.1
                                                                                                                          TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: SITE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LOCATION: (3)
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; Publication No. US2003000355A1
; GENERAL INFORMATION:
; APPLICANT: ROSEN et al.
; TITLE OF INVENTION: 90 Human Secreted Proteins
; FILE REFRENCE: PZ013P1
; CURRENT APPLICATION WHBER: US/09/774,639
; CURRENT PILING DATE: 2001-07-09
; PRIOR APPLICATION NUMBER: BARLIER APPLICATION NUMBER: 09/244,112
; PRIOR FILING DATE: EARLIER FILING DATE: 1999-02-04
; NUMBER OF SEQ ID NOS: 371
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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CURRENT FILING DATE: 1999-05-27
EARLIER APPLICATION NUMBER: 60/087,340
EARLIER FILING DATE: 1998-05-29
EARLIER FILING DATE: 1998-05-29
EARLIER FILING DATE: 1998-09-10
EARLIER FILING DATE: 1998-09-10
EARLIER FILING DATE: 1999-04-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Reinhard Ebner
APPLICANT: Steven M. Ruben
TITLE OF INSTRIDON: INTERLEUKINS-21 AND 22
FILE REFERENCE: PF470
CURRENT FILING DATE: 2002-03-01
PRICA APPLICATION NUMBER: PCT/GB00/03287
PRIOR PILING DATE: 2000-08-24
PRIOR APPLICATION NUMBER: 9920673.2
PRIOR FILING DATE: 1999-09-01
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FASTSEQ FOR Windows Version 4.0
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Publication No. US20030003545A1
GENERAL INFORMATION:
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: arthropod
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US-09-774-639-2
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TITLE OF INVENTION: 31 Human secreted proteins
FILE REFERENCE: P2026P1
CURRENT APPLICATION NUMBER: US/09/798,889
CURRENT FILING DATE: 2001-03-06
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 09/393,022
PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/077,714
PRIOR APPLICATION NUMBER: EARLIER PELICATION NUMBER: 60/077,714
PRIOR PILING DATE: BARLIER FILING DATE: 1998-03-12
PRIOR PILING DATE: BARLIER FILING DATE: 1998-03-12
PRIOR PILING DATE: EARLIER FILING DATE: 1998-03-12
PRIOR PILING DATE: BARLIER FILING DATE: 1998-03-12
PRIOR FILING DATE: BARLIER FILING DATE: 1998-03-12
NUMBER OF SEQ ID NOS: 185
SOUTHWARE: PARCHIER FILING DATE: 1998-03-12
SOUTHWARE: PARCHIER FILING DATE: 1998-03-12
LENGTH: 5
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Publication No. US20030009018A1
GENERAL INFORMATION:
APPLICANT: Maeda, Masatsugu
APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERBUCE: 06501-1051031
CURRENT APPLICATION NUMBER: US/10/105,930
                                                                                                                                                                                                                                                                                                                      Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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SOFTWARE: Patentin Ver. 2.0
                    SEQ ID NO 2
LENGTH: 5
TYPE: PRT
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: Site
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APPLICANT: Waeda, Masaleugu

APPLICANT: Yaguchi, No. US2003009018Aliko

TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12

FILE REPERENCE: 06501-105US1

CURRENT APPLICATION NUMBER: US/10/105,930

CURRENT APPLICATION NUMBER: PCT/JP00/06654

PRIOR PRIOR PELICANION NUMBER: D200-09-27

PRIOR PELING DATE: 2000-09-27

PRIOR PELING DATE: 1999-09-27

PRIOR FILING DATE: 1999-09-27

PRIOR FILING DATE: 1999-09-27

NUMBER OF SEQ ID NOS: 77

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 59
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APPLICANT: Maeda, Masatsugu
APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-1051031
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: JP 2000-240397
                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 9; Le
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0;
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: JP 2000-240397
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 1000-09-03
PRIOR FILING DATE: 1099-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FASTSEQ for Windows Version 4.0
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; Sequence 59, Application US/10105930
; Publication No. US2003009918A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                     ) ORGANISM: Homo sapiens
US-10-105-930-58
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US-10-105-930-59
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US-10-105-930-60
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APPLICAUT: Maeda, Masatsugu
APPLICAUT: Maeda, Masatsugu
APPLICAUT: Maeda, Mosu US2003009018Aliko
ITILE OP INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 66501-105451
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: JP 2000-240397
PRIOR APPLICATION NUMBER: JP 11-273358
PRIOR PLING DATE: 2000-08-03
PRIOR FILING DATE: 1999-09-77
NUMBER OF SEQ ID NOS: 77
SOFTWARE PAECES FOR WINDOWS Version 4.0
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Publication No. US2003009018A1
GENERAL INFORMATION:
APPLICANT: Maeda, Masatsugu
APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOISTIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-105US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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                  PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FREUESEQ for Windows Version 4.0
SEQ ID NO 21
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                          ) LOCATION: 3
; OTHER INFORMATION: Xaa = any amino acid US-10-105-930-21
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                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: Exemplary motif
        CURRENT FILING DATE: 2002-03-25
                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-105-930-57
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LENGTH: 5
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PUBLICATION NO. US20030009018A1

GENERAL INFORMATION:
APPLICANT: Maeda, Masatsugu
APPLICANT: Vaguchi, No. US2003009018A1iko
ITILE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 66501.105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR PELING DATE: 2000-09-27
PRIOR PELING DATE: 2000-09-27
PRIOR PELING DATE: 2000-09-80
PRIOR PELING DATE: 2000-09-7
PRIOR PELING DATE: 2000-09-7
PRIOR PELING DATE: 2000-09-7

PRIOR APPLICATION NUMBER: UP 11-273358

PRIOR PILING DATE: 1999-09-27

NUMBER OF SEQ ID NOS: 77

SOFTWARE FERSE FASTSEQ for Windows Version 4.0
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APPLICANT: Maeda, Masatsugu
APPLICANT: Maeda, Masatsugu
APPLICANT: Maeda, Masatsugu
APPLICANT: Vaguchi, No. US2003009018Aliko
TITLE OF INVENTION: NOVEL HEMOPOLETIN RECEPTOR PROTEIN, NR12
FILE REPERENCE: 65501-105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: JP 2000-240397
PRIOR APPLICATION NUMBER: JP 11-273358
PRIOR FILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOPTWARE: FASTSEQ for Windows Version 4.0
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PRIOR FILING DATE: 2000-08-03
PRIOR APPLICATION NUMBER: JP 11-273358
PRIOR FILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 60
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                                                                                                                                                                                          TYPE: PRT
ORGANISM: Homo sapiens
US-10-105-930-60
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Matches 1; Conserva
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LENGTH: 5
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| Sequence 63, Application US/10105930
| Publication No. US20030009018A1
| GENERAL INFORMATION:
| APPLICANT: Macda, Masatsugu
| APPLICANT: Yaquchi, No. US20030009018A11kc
| TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
| FILE REFERENCE: 06504-105U81
| CURRENT PAPLICATION NUMBER: US/10/105,930
| CURRENT PAPLICATION NUMBER: PCT/0F00/06654
| PRIOR APPLICATION NUMBER: PCT/0F00/06654
| PRIOR PLING DATE: 2000-09-27
| PRIOR PLING DATE: 2000-09-27
| PRIOR PLING DATE: 1900-09-27
| PRIOR PLING DATE: 1999-09-27
| NUMBER OF SEQ ID NOS: 77
| SOFTWARE FastSEQ for Windows Version 4.0
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Publication No. US20030009018A1

FORDERAL INFORMATION:
APPLICANT: Maeda, Masatsugu
APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-105031
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
FRIOR RAPLICATION NUMBER: PCT/JD00/06654
FRIOR PAPLICATION NUMBER: PT 2000-240397
FRIOR PILING DATE: 2000-08-03
FRIOR PILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE FESTSEQ for Windows Version 4.0
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                                                                                                                                                      Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
; SEQ ID NO 62
; LENGTH: 5
; TYPE: PRT
; ORGANISM: Homo sapiens
US-10-105-930-62
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LENGTH: 5
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US-10-105-390-68
Sequence 68, Application US/10105930
Sequence 68, Application US/10105930
Publication No. US20030009018A1
Sequence 10 No. US20030009018A1
SERENAL INFORMATION:
TITLE OF INVENTION: NOVEL HEMOPOLETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-105U31
CURRENT PLING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR PILING DATE: 2000-09-27
PRIOR FILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFFWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                            APPLICANT: Maeda, Masateugu
APPLICANT: Maeda, Masateugu
APPLICANT: Maeda, Masateugu
APPLICANT: Yaguchi, No. US20030009018Aliko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERENCE: 06501-105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR FILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 1999-09-27
PRIOR PELING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 67
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; Sequence 67, Application US/10105930
; Publication No. US20030009018A1
; GENERAL INFORMATION:
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US-10-105-930-68
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CORGANISM: Homo sapiens
US-10-105-930-67
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GENERAL INFORMATION:
APPLICANT: Masatsugu
FAPPLICANT: Masatsugu
FILE REPRENCE: 06501-105403
CURRENT APPLICATION NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-1054031
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 1999-09-27
PRIOR FILING DATE: 1999-09-27
PRIOR FILING DATE: 1999-09-27
SPRIOR FILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastESQ for Windows Version 4.0
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APPLICANT: Meaateugu
APPLICANT: Maguchi, No. US20030009018Aliko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 66501-105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25;
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 1999-09-09-7
NUMBER OF SEQ ID NOS: 77
SOFTWARE: FastSEQ for Windows Version 4.0
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                    47.8%; Score 11; DB 9; Length 5; 100.0%; Pred. No. 1.2e+05; tive 0; Mismatches 0; Indels
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US-10-105-930-66
Sequence 66, Application US/10105930
Publication No. US20030009018A1
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ORGANISM: Homo sapiens
US-10-105-930-66
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US-10-105-930-65
                    Query Match
Best Local Similarity
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US-10-105-930-65
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APPLICANT: Masat sugu
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APPLICANT: Masat sugu
APPLICANT: Maguchi, No. US20030009018A1iko
TITLE OP INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERBNCE: 06501-105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: JP 2000-240397
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
PRIOR PILING DATE: 2000-08-03
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PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 1000-08-03
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APPLICANT: Maeda, Masatsugu
APPLICANT: Yaguchi, No. US20030009018Aliko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERENCE: 055501-1050131
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: JP 2000-240397
PRIOR APPLICATION NUMBER: JP 2000-240397
PRIOR PILING DATE: 1999-09-27
PRIOR PILING DATE: 1999-09-27
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SOFTWARE: FastSEQ for Windows Version 4.0
                                                                  Sequence 69, Application US/10105930 Publication No. US20030009018A1 GENERAL INFORMATION:
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US-10-105-930-72
'Sequence 72, Application US/10105930
'Publication No. US20030009018A1
'GENERAL INFORMATION:
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US-10-105-930-69
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US-10-105-930-71
RESULT 61
US-10-105-930-69
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APPLICANT: Maeda, Masatsugu
APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERENCE: 66501-105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT APPLICATION NUMBER: US/10/10654
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 2000-08-03
PRIOR FILING DATE: 1999-09-27
NUMBER: PSEQ ID NOS: 77
SOPTWARE: FastSEQ for Windows Version 4.0
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APPLICANT: Masatsugu
APPLICANT: Masatsugu
APPLICANT: Masatsugu
APPLICANT: Yaguchi, No. US20030009018Aliko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06554
PRIOR APPLICATION NUMBER: DCT/JP00/06554
PRIOR APPLICATION NUMBER: D200-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
PRIOR PILING DATE: 2000-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE PRICES FASISEQ for Windows Version 4.0
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APPLICANT: Yaguchi, No. US20030009018A11ko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REPERENCE: 065501-1059031
CURRENT APPLICATION NUMBER: US/10/105,930
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47.8%; Score 11; DB 9; Length 5;
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 73, Application US/10105930 ; Publication No. US20030009018A1 ; GENERAL INFORMATION:
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US-10-105-930-72
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PRIOR APPLICATION NUMBER: JP 11-273358
PRIOR FILING DAFF: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOFTWARE: PASTSEQ FOR Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Unknown US-09-262-126C-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        , APPLICANT: Miller, Brian S.
                                                                                                                         TYPE: PRT
CRGANISM: Homo sapiens
US-10-105-930-76
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                                                                                          SEQ ID NO 76
LENGTH: 5
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LENGTH: 5
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JETOTOMATE: Maeda, Masateugu
APPLICANT: Yaguchi, No. US2003009918Aliko
JITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILER REFERENCE: 06501-105USI
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR APPLICATION NUMBER: DP 2000-240397
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 1999-09-27
NUMBER OF SEQ ID NOS: 77
SOSTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 75
LENGTH: 5
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APPLICANT: Yaguchi, No. US20030009018A1iko
TITLE OF INVENTION: NOVEL HEMOPOIETIN RECEPTOR PROTEIN, NR12
FILE REFERENCE: 06501-105US1
CURRENT APPLICATION NUMBER: US/10/105,930
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR PILING DATE: 2000-09-03
PRIOR FILING DATE: 2000-08-03
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47.8%; Score 11; DB 9; Le
Best Local Similarity 100.0%; Pred. No. 1.2e+05;
Matches 1; Conservative 0; Mismatches 0;
CURRENT FILING DATE: 2002-03-25
PRIOR APPLICATION NUMBER: PCT/JP00/06654
PRIOR FILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: JP 2000-240397
PRIOR FILING DATE: 2000-08-03
PRIOR PILING DATE: 1999-09-27
PRIOR PILING DATE: 1999-09-27
SOFTWARE: FASTSEQ for Windows Version 4.0
SEQ ID NO 74
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Sequence 76, Application US/10105930
Publication No. US20030009018A1
GENERAL INFORMATION:
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; ORGANISM: Homo sapiens
US-10-105-930-74
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US-10-105-930-75
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Best Local Similarity
Matches 1; Conserv
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ITILE OF INVENTION: Nucleic Acids, Proteins, and Antibodies FILE REPERENCE: PUZOB
CURRENT APPLICATION UNDBER: US/09/989,442
CURRENT FILING DATE: 2001-11-21
PRIOR PPLICATION NUMBER: 60/179,065
PRIOR PPLICATION NUMBER: 60/180,628
PRIOR PLING DATE: 2000-02-04
PRIOR APPLICATION NUMBER: 60/214,886
PRIOR APPLICATION NUMBER: 60/214,886
PRIOR APPLICATION NUMBER: 60/217,487
PRIOR APPLICATION NUMBER: 60/217,487
PRIOR APPLICATION NUMBER: 60/225,758
PRIOR APPLICATION NUMBER: 60/225,758
PRIOR APPLICATION NUMBER: 60/225,758
PRIOR PILING DATE: 2000-08-14
PRIOR PLING DATE: 2000-08-14
PRIOR PLING DATE: 2000-07-25
PRIOR FILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-07-21
Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Shetty, Jayarama K.
TITLE OF INVENTION: Modified Forms of Pullulanase
FILE REPERENCE: GC396-2
CURRENT APPLICATION NUMBER: US/09/262,126C
CURRENT FILING DATE: 1999-03-03
NUMBER OF SEQ ID NOS: 9
SOFTWARE: FastSEQ for Windows Version 3.0
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R FILING DATE: 2000-08-14

R APPLICATION NUMBER: 60/226,868

R FILING DATE: 2000-08-22

R APPLICATION NUMBER: 60/216,647

R FILING DATE: 2000-07-07

R APPLICATION NUMBER: 60/225,267 FILING DATE: 2000-09-05
APPLICATION NUMBER: 60/236,367
FILING DATE: 2000-09-29
APPLICATION NUMBER: 60/237,039
FILING DATE: 2000-10-02 APPLICATION NUMBER: 60/236,327 FILING DATE: 2000-09-29 APPLICATION NUMBER: 60/241,785 ALEING DATE: 2000-10-20 APPLICATION NUMBER: 60/244,617 APPLICATION NUMBER: 60/251,868 FILING DATE: 2000-12-08 APPLICATION NUMBER: 60/229,344 FILING DATE: 2000-09-01 APPLICATION NUMBER: 60/234,997 APPLICATION NUMBER: 60/229,513
FILING DATE: 2000-09-01
FILING DATE: 2000-09-05
APPLICATION NUMBER: 60/231,413
FILING DATE: 2000-09-08 APPLICATION NUMBER: 60/218,290 FILING DATE: 2000-07-14 APPLICATION NUMBER: 60/225,757 APPLICATION NUMBER: 60/241,809 FILING DATE: 2000-10-20 APPLICATION NUMBER: 60/249,299 APPLICATION NUMBER: 60/229,343 FILING DATE: 2000-09-01 APPLICATION NUMBER: 60/229,345 FILING DATE: 2000-07-07 APPLICATION NUMBER: 60/225,270 FILING DATE: 2000-08-14 APPLICATION NUMBER: 60/251,869 FILING DATE: 2000-12-08 FILING DATE: 2000-09-27 APPLICATION NUMBER: 60/234,274 APPLICATION NUMBER: 60/234,223 FILING DATE: 2000-09-21 APPLICATION NUMBER: 60/228,924 FILING DATE: 2000-08-30 APPLICATION NUMBER: 60/224,518 FILING DATE: 2000-08-14 APPLICATION NUMBER: 60/236,369 FILING DATE: 2000-09-29 APPLICATION NUMBER: 60/220,964 FILING DATE: 2000-07-26 APPLICATION NUMBER: 60/225,268 60/236,368 FILING DATE: 2000-09-29 APPLICATION NUMBER: 60/251,856 FILING DATE: 2000-09-01 APPLICATION NUMBER: 60/229,287 FILING DATE: 2000-09-08 APPLICATION NUMBER: 60/229,509 FILING DATE: 2000-10-02 APPLICATION NUMBER: 60/237,038 APPLICATION NUMBER: 60/225,447 APPLICATION NUMBER: 60/216,880 APPLICATION NUMBER: 60/224,519 APPLICATION NUMBER: 60/235,834 FILING DATE: 2000-08-14 FILING DATE: 2000-12-08 FILING DATE: 2000-08-14 FILING DATE: 2000-09-21 FILING DATE: 2000-08-14 FILING DATE: 2000-11-17 FILING DATE: 2000-11-01 FILING DATE: 2000-09-25 APPLICATION NUMBER: PRIOR PRIOR
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REFLIGATION NUMBER: 60/237,040
REPLING DATE: 2000-10-02
REFLING DATE: 2000-10-02
REFLING DATE: 2000-10-02
REFLING DATE: 2000-10-20
REFLING DATE: 2000-10-20
REPLING DATE: 2000-10-20
REPLING DATE: 2000-10-20 FILING DATE: 2000-09-14 APPLICATION NUMBER: 60/231,242 FILING DATE: 2000-09-08 APPLICATION NUMBER: 60/227,182 FILING DATE: 2000-08-22 APPLICATION NUMBER: 60/225,214 FILING DATE: 2000-10-02 APPLICATION NUMBER: 60/236,370 FILING DATE: 2000-09-29 FILING DATE: 2000-10-13
APPLICATION WUMBER: 60/239,937
APPLICATION DATE: 2000-10-13
APPLICATION NUMBER: 60/241,787 FILING DATE: 2000-11-08
APPLICATION NUMBER: 60/246,532
FILING DATE: 2000-11-08 FILING DATE: 2000-06-30 APPLICATION NUMBER: 60/225,266 FILING DATE: 2000-08-14 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,207 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,245 FILING DATE: 2000-11-17 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,297 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/232,400 APPLICATION NUMBER: 60/236,802 FILING DATE: 2000-10-02 FILING DATE: 2000-10-20 APPLICATION NUMBER: 60/246,474 APPLICATION NUMBER: 60/249,216 FILING DATE: 2000-11-17
APPLICATION NUMBER: 60/249,210 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/226,681 FILING DATE: 2000-08-22 APPLICATION NUMBER: 60/225,759 APPLICATION NUMBER: 60/225,213 LICATION NUMBER: 60/235,836 APPLICATION NUMBER: 60/230,438 FILING DATE: 2000-09-06
APPLICATION NUMBER: 60/215,135 APPLICATION NUMBER: 60/249,218 APPLICATION NUMBER: 60/249,208 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,213 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,212 APPLICATION NUMBER: 60/249,244 APPLICATION NUMBER: 60/249,217 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,211 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,215 APPLICATION NUMBER: 60/249,264 FILING DATE: 2000-11-17 APPLICATION NUMBER: 60/249,214 APPLICATION NUMBER: 60/237,037 FILING DATE: 2000-09-27 FILING DATE: 2000-08-14 2000-08-14 FILING DATE: 2000-08-14 FILING DATE: 2000-11-17 2000-11-17 FILING DATE: FILING DATE: FILING DATE: PRIOR PRIOR

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GREATURE TILING DATE: 1998-02-06

TILE OF INVENTION: Dendritic Enriched Secreted Lymphocyte Activation TILE OF INVENTION: Dendritic Enriched Secreted Lymphocyte Activation TILE OF INVENTION: Dendritic Enriched Secreted Lymphocyte Activation TILE OF INVENTION: Molecule
FILE REFERENCE: PF448
CURRENT PELLING NAMBER: US/09/369,248
CURRENT FILING DATE: 1999-08-05
EARLIER FILING DATE: 1998-0-06
EARLIER FILING DATE: 1998-0-06
EARLIER FILING DATE: 1998-0-06
SAFLIER FILING DATE: 1998-0-06
SOFTWARE: PATENTIN VET: 2.0
SOFTWARE: PATENTIN VET: 2.0
SEQ ID NO 5
LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 47.8%; Score 11; DB 9; Length 5; Best Local Similarity 100.0%; Pred. No. 1.2e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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TITLE OF INVENTION: Human Neuropeptide Receptor
FILE REFERENCE: PF168P2
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; OTHER INFORMATION: Xaa equals any amino acid
US-09-393-696-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; LOCATION: (3)
; OTHER INFORMATION: Xaa equals any amino acid US-09-369-248-5
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CURRENT FILING DATE: 1990-09-10
EARLIER APPLICATION NUMBER: ECT/US95/05616
EARLIER FILING DATE: 1995-05-05
EARLIER FILING DATE: 1995-06-05
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver: 2.0
SEQ ID NO 14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; Sequence 14, Application US/09393696
; Publication No. US20030022277A1
; GENERAL INFORMATION:
                                                                                                                                                                               US-09-369-248-5
; Sequence 5, Application US/09369248
; Publication No. US20030022276Al
; GENERAL INFORMATION:
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US-09-393-696-14
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FEATURE:
OTHER HYPORMATION: alpha5betal integrin-binding peptide
US-10-116-067-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: LINDBOM, Lennart
APPLICANT: LINDBOM, Lennart
APPLICANT: WERR, Joachim
TITLE OF INVENTION: Anti-Inflammatory Medicaments
FILE REPERBNCE: 1291-0200P
CURRENT APPLICATION NUMBER: US/10/116,067
CURRENT PILING DATE: 2002-04-05
PRIOR FILING DATE: 1998-10-07
PRIOR FILING DATE: 1998-10-07
PRIOR APPLICATION NUMBER: US 60/103,558
PRIOR PILING DATE: 1999-10-07
                      PRIOR FILING DATE: 2000-09-08

PRIOR APPLICATION NUMBER: 60/232,080

PRIOR PILING DATE: 2000-09-08

PRIOR PILING DATE: 2000-09-08

PRIOR PELING DATE: 2000-09-08

PRIOR PILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR FILING DATE: 2000-09-08

PRIOR PILING DATE: 2000-09-14

PRIOR PELING DATE: 2000-09-14

PRIOR PILING DATE: 2000-10-20

PRIOR PILING DATE: 2000-10-8

PRIOR PILING DATE: 2000-10-33,065
APPLICATION NUMBER: 60/232,081
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Publication No. US20030012780A1
GENERAL INFORMATION:
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FEATURE:
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PRIOR APPLICATION NUMBER: BARLIER APPLICATION NUMBER: 09/227,357

PRIOR PLIING DATE: EARLIER FILING DATE: 1990-0.08

PRIOR APPLICATION NUMBER: BARLIER PLING DATE: 1996-07-07

PRIOR PLIING DATE: EARLIER FILING DATE: 1996-07-07

PRIOR PLIING DATE: EARLIER FILING DATE: 1996-07-07

PRIOR PLIING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR PPLICATION NUMBER: EARLIER PLIING DATE: 1997-07-08

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PRIOR FILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR FILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR FILING DATE: EARLIER PLIING DATE: 1997-07-09

PRIOR FILING DATE: EARLIER PLIING DATE: 1997-07-09
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TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR BREAST CANCER
FILE REPERENCE: A-69959/RMS/DCF
CURRENT APPLICATION NUMBER: US/09/747,377
CURRENT FILING DATE: 2000-12-22
NUMBER OF SEQ ID NOS: 493
SOFTWARE: Patentin version 3.0
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TITLE OF INVENTION: 123 Human Secreted Proteins
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CURRENT APPLICATION NUMBER: US/09/983,802
CURRENT FILING DATE: 2001-10-25
                                                                                                                                                                                  Sequence 493, Application US/09747377
Publication No. US20030022255A1
GENERAL INFORMATION:
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Publication No. US20030022185A1
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GENERAL INFORMATION:
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Best Local Similarity
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OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino OTHER INFORMATION: acids
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TITLE OF INVENTION: Dendritic Enriched Secreted Lymphocyte Activation
TITLE OF INVENTION: Molecule
FILE REFERENCE: PP448P2
CURRENT APPLICATION NUMBER: US/10/062,523
PRIOR FILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR PILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR PILING DATE: EARLIER FILING DATE: 1997-07-08

PRIOR FILING DATE: EARLIER FILING DATE: 1997-07-08-18

PRIOR FILING DATE: EARLIER FILING DATE: 1997-07-08-18

PRIOR FILING DATE: EARLIER FILING DATE: 1997-07-08-18

PRIOR FILING DATE: EARLIER FILING DATE: 1997-08-18

PRIOR FILING DATE: EARLIER FILING DATE: 1997-08-18

PRIOR FILING DATE: EARLIER FILING DATE: 1997-08-18

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/055,949

PRIOR FILING DATE: EARLIER FILING DATE: 1997-08-18

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/055,949

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/055,949

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/055,949

PRIOR APPLICATION NUMBER: EARLIER APPLICATION NUMBER: 60/055,944

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PRIOR APPLICATION NUMBER: EARLIER PILING DATE: 1997-08-18

PRIOR APPLICATI
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SOFTWARE: Patentin Ver. 2.0
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US-10-062-523-5
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LENGTH: 5
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CURRENT FILING DATE: 2002-02-05

PRIOR PAPLICATION NUMBER: 60/267,523

PRIOR PILING DATE: 2000-02-06

PRIOR PILING DATE: 2000-02-06

PRIOR PILING DATE: 2000-03-03

PRIOR PILING DATE: 1000-03-03

PRIOR PILING DATE: 1000-03-17

PRIOR PILING DATE: 1090-03-17

PRIOR PILING DATE: 1090-06-05

PRIOR PILING DATE: 1090-06-05

PRIOR PILING DATE: 1090-06-05

PRIOR PILING DATE: 1090-06-05

PRIOR PILING DATE: 1090-02-04

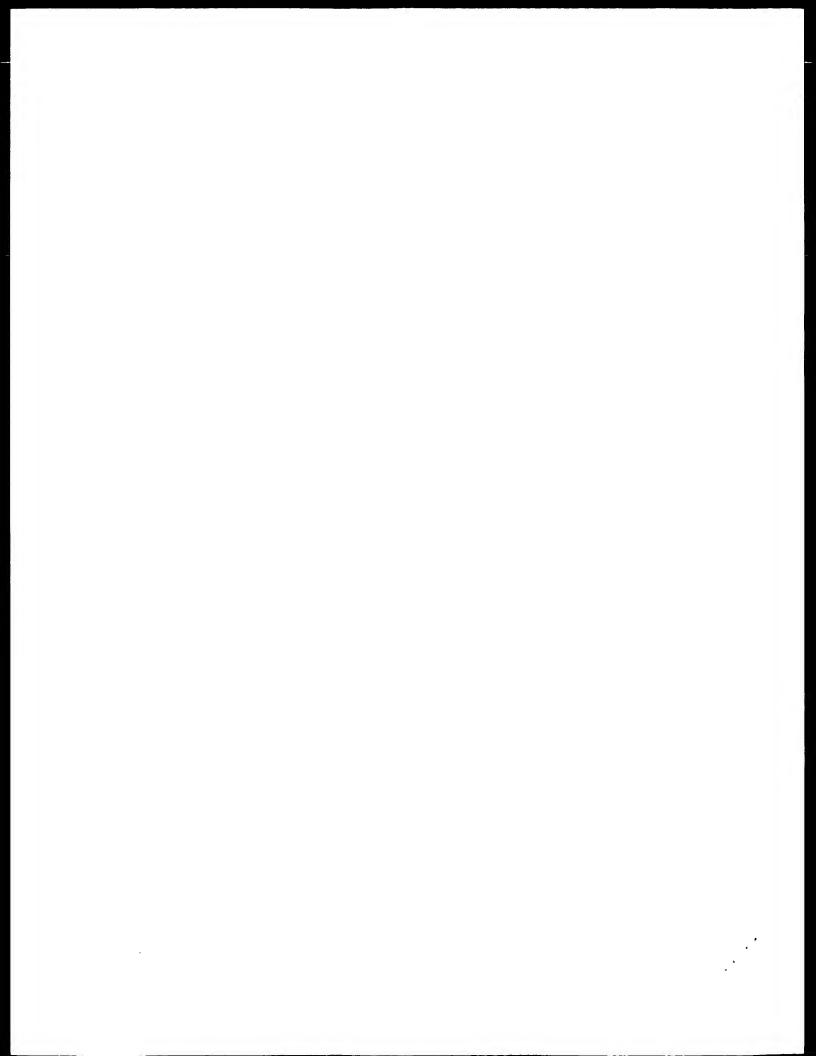
PRIOR PILING DATE: 1090-03-17

PRIOR PILING DATE: 1090-03-17

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Sequence 2, Appli Sequence 3, Appli Sequence 3, Appli Sequence 32, Appli Sequence 791, App Sequence 139, App Sequence 139, Appli Sequence 10, Appli Sequence 10, Appli Sequence 11, Appli

Sequence 11, Appl Sequence 11, Appl Sequence 13, Appl Sequence 13, Appl Sequence 11, Appl Sequence 11, Appl Sequence 11, Appl Sequence 11, Appl Sequence 100, App Sequence 228, App Sequence 14, Appl Sequence 15, Appl

Sequence:

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Searched:

Database

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6 US-09-214-371-2

6 US-09-214-371-3

18 US-09-403-440A-1

18 US-09-403-440A-1

18 US-09-403-440A-1

18 US-09-403-440A-1

18 US-09-957-806A-226

18 US-09-957-806A-226

18 US-09-357-10

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18 US-09-351-955A-16-791

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18 US-09-515-955A-16-791

18 US-09-515-955A-16-791

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18 US-09-515-955A-19-9

18 US-09-5
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US-10-197-954-139
US-10-200-923-8
US-10-200-923-10
PCT-US01-20969-3
US-09-214-371-2
US-09-281-717-3
US-09-403-440A-1
   Sequence 11, Appl
Sequence 11, App
Sequence 11, Appl
Sequence 114, App
Sequence 114, Appl
Sequence 111, Appl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     the number of results predicted by chance to have a r than or equal to the score of the result being printed, ed by analysis of the total score distribution.
                                                                                  February 12, 2003, 10:51:44; Search time 131 Seconds (without alignments) 24.608 Million cell updates/sec
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1: \cgn2 6/ptodata/1/paa/PCTUS COMB.pep:*

2: \cgn2 6/ptodata/1/paa/PCTUS COMB.pep:*

3: \cgn2 6/ptodata/1/paa/US08 COMB.pep:*

4: \cgn2 6/ptodata/1/paa/US08 COMB.pep:*

5: \cgn2 6/ptodata/1/paa/US08 COMB.pep:*

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7: \cgn2 6/ptodata/1/paa/US08 COMB.pep:*

8: \cgn2 6/ptodata/1/paa/US08 COMB.pep:*

10: \cgn2 6/ptodata/1/paa/US08 COMB.pep:*

11: \cgn2 6/ptodata/1/paa/US08 COMB.pep:*

12: \cgn2 6/ptodata/1/paa/US08 COMB.pep:*

13: \cgn2 6/ptodata/1/paa/US08 COMB.pep:*

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15: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

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17: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

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19: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

20: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

21: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

22: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

23: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

24: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

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27: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

28: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

29: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

20: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

22: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

23: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*

24: \cgn2 6/ptodata/1/paa/US09 COMB.pep:*
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               GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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PCT-US99-26277-114
US-09-601-558-11
US-09-811-047B-114
US-09-891-047C-114
US-09-894-594-11
                                                                                                                                                                                                                                          hits satisfying chosen parameters:
                                                                                                                                                                                                                    4569144 seqs, 644733110 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             6/ptodata/1/paa/US60
                                                                                                                                                                                                                                                                                                    Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 100 summaries
                                                            - protein search, using sw model
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36, Appl 95, Appl 139, App 8, Appli

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Sequence Sequence

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80 15 65.2 5 26 US-10-200-923-11 Sequence 11, Appl 81 15 65.2 5 26 US-10-200-923-12 Sequence 12, Appl 83 15 65.2 5 26 US-10-200-923-13 Sequence 12, Appl 84 15 65.2 5 26 US-10-230-933-13 Sequence 11, Appl 85 14 60.9 5 17 US-09-392-9418-99 Sequence 65, Appl 86 14 60.9 5 17 US-09-392-9418-99 Sequence 65, Appl 87 14 60.9 5 17 US-09-392-9418-19 Sequence 81, Appl 88 14 60.9 5 17 US-09-392-9418-19 Sequence 67, Appl 89 13 56.5 5 1 PCT-US00-07677-97 Sequence 7, Appl 90 13 56.5 5 1 PCT-US02-1448-4 Sequence 7, Appl 91 13 56.5 5 1 PCT-US02-1448-4 Sequence 7, Appl 92 13 56.5 5 1 PCT-US02-1448-4 Sequence 7, Appl 94 13 56.5 5 1 PCT-US02-1448-4 Sequence 11, Appl 95 13 56.5 5 3 US-07-69-9256-11 Sequence 17, Appl 96 13 56.5 5 3 US-07-69-755-47 Sequence 47, Appl 97 13 56.5 5 3 US-07-69-755-47 Sequence 49, Appl 97 13 56.5 5 3 US-07-69-755-47 Sequence 49, Appl 97 13 56.5 5 3 US-07-69-755-47 Sequence 49, Appl 98 13 56.5 5 3 US-07-69-755-49 Sequence 47, Appl 99 13 56.5 5 3 US-07-69-755-49 Sequence 49, Appl 99 13 56.5 5 3 US-07-69-755-49 Sequence 47, Appl 99 13 56.5 5 3 US-07-69-755-49 Sequence 47, Appl 99 13 56.5 5 3 US-07-69-755-49 Sequence 47, Appl 99 13 56.5 5 3 US-07-69-755-49 Sequence 47, Appl 99 13 56.5 5 3 US-07-69-755-49 Sequence 47, Appl 99 13 56.5 5 3 US-07-69-755-49 Sequence 47, Appl 99 13 56.5 5 3 US-07-89-08-47 Sequence 47, Appl 99 13 56.5 5 3 US-07-89-08-47 Sequence 47, Appl 99 13 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 13 56.5 5 3 US-07-89-08-47 Sequence 47, Appl 99 13 56.5 5 3 US-07-89-08-47 Sequence 47, Appl 99 13 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-89-755-49 Sequence 47, Appl 99 14 56.5 5 3 US-07-8
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PCT-US99-02405-11
Sequence 11, Application PC/TUS9902405
GENERAL INCOMETATION:
TITLE OF INVENTION: Peptide Mimotopes of Carbohydrate Antigens
TITLE OF INVENTION DATE: 1999-02-04
MUMMERS OF SEQ ID NOS: 60
SEQ ID NO: 60
S
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Wistar Institute of the University of Pennsylvania
APPLICANT: Blaszczyk-Thurin, Magdalena
APPLICANT: Rieber-Emmons, Thomas
APPLICANT: Rieber-Emmons, Thomas
TITLE OF INVENTION: Compositions and Methods For Treatment of Cancer FILE REFERENCE: WST93AUSA
CURRENT APPLICATION NUMBER: 2001-08-14
PRIOR APPLICATION NUMBER: 60/107,478
PRIOR APPLICATION NUMBER: 60/107,478
PRIOR APPLICATION NUMBER: 60/107,478
SEQ ID NOS: 121
SOFTWARE: PatentIn Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ## Sequence 11, Application US/09601558
## GENERAL INFORMATION:
## GENERAL INFORMATION:
## APPLICANT: Kieber-Emmons, Thomas
## APPLICANT: The Trustees of the University of Pennsylvania
## TITLE OF INVENTION: Peptide Mimotopes Of Carbohydrate Antigens
## FILE REFERENCE: UR13567
## CURRENT APPLICATION NUMBER: US/09/601,558
## CURRENT PILING DATE: 2000-08-03
## PRIOR APPLICATION NUMBER: 60/073,690
## PRIOR PILING DATE: 1999-02-04
## PRIOR PILING DATE: 1999-02-04
## PRIOR PILING DATE: 1998-02-04
## PRIOR PILING DATE: 1998-02-04
## NUMBER OF SEQ ID NOS: 60
## SOFTWARE: PatentIn Ver. 2.1
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                                                                                                                                                                                                                       87.0%; Score 20; DB 1; Length 5; 60.0%; Pred. No. 4.2e+06; Live 0; Mismatches 2; Indels
                                                                                                             OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: peptido-mimetic of a Lewis antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Description of Artificial Sequence: OTHER INFORMATION: peptido-mimetic of a Lewis antigen
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-831-047B-114; Sequence 114, Application US/09831047B; GENERAL INFORMATION:
                           TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                          Query Match
Best Local Similarity 60.0°
Matches 3; Conservative
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Best Local Similarity 60.09
Matches 3; Conservative
                                                                                                                                                 ; OTHER INFORMALL PCT-US99-26277-114
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US-09-601-558-11
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LENGTH: 5
LENGIH: 5
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Gaps

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TITLE OF INVENTION: METHODS AND COMPOUNDS FOR MODULATING NUCLEAR RECEPTOR TITLE OF INVENTION: ACTIVITY ACTIVITY

TITLE OF INVENTION: ACTIVITY
FILE REPRENCE: 9811-016-228

CURRENT APPLICATION NUMBER: PCT/US01/20969

CURRENT FILING DATE: 2000-06-30

PRIOR APPLICATION NUMBER: US 60/079,965

PRIOR PILING DATE: 1998-03-30

PRIOR PILING DATE: 1998-03-30

PRIOR PILING DATE: 1998-03-30

PRIOR PILING DATE: 1999-03-30

PRIOR PILING DATE: 1999-03-30

PRIOR PILING DATE: 1999-03-30

NUMBER OF SEQ ID NOS: 83

SOFTWARE: PATENTIN VEXESION 3.0
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          87.0%; Score 20; DB 22; Length 5; 60.0%; Pred. No. 4.2e+06; 1ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
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OTHER INFORMATION: Xaa is any amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Chene, Patrick
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the I
                                                                                                                                                                                                                                                                          GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: GUY, RODNEY
APPLICANT: DAKTREW, JOHN
APPLICANT: DAKTREW, JOHN
APPLICANT: PENG, WELJUN
APPLICANT: PENG, WELJUN
APPLICANT: PETER, KUSHNER
APPLICANT: PETER, KUSHNER
APPLICANT: PETER, WEST
APPLICANT: ANAMOTO, WEBT
APPLICANT: YARAMOTO, KETTH
APPLICANT: YARAMOTO, KETTH
APPLICANT: YARAMOTO, KETTH
APPLICANT: KUNIZ, IRWIN
APPLICANT: KUNIZ, IRWIN
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; APPLICANT: Lane, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 100.
Matches 5; Conservative
                                                            3; Conservative
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                                      Best Local Similarity
                                                                                                             1 FXXLW 5
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US-09-214-371-2
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APPLICANT:
               Query Match
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APPLICANT: Kieber-Emmons, Thomas
APPLICANT: Weiner, David B.
APPLICANT: Weiner, David B.
APPLICANT: Weiner, David B.
APPLICANT: Weiner, David B.
APPLICANT: Wonzard: Karbassi, Behjatolah
ITILE OP INVENTION: Same
FILE OP INVENTION: Same
FILE OF INVENTION: Same
FILE OF INVENTION: Same
FILE OP INVENTION NUMBER: US/09/894,594
FILE OF TARENT FILING DATE: 2000-11-07
FILE OP INVENTION NUMBER: E0/073,690
FILE OF FILE OP INVENTION NUMBER: E0/073,690
FILE OF FILE OP INVENTION NUMBER: E0/214,517
FILE OF SEQ ID NOS: 70
SOFTMARE: Patent In version 3.0
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                                                                                                                                                                                                                                                                                                                   Sequence 114, Application US/09831047C
; GENERAL INFORMATION:
APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Trustees of the University of Pennsylvania
APPLICANT: Blaszczyk-Thurin, Magdalena
APPLICANT: Kieber-Emmons, Thomas
TILLE OF INVENTION: Compositions and Methods For Treatment of Cancer
FILE REFERENCE: WST93ANGSA
CURRENT APPLICATION NUMBER: US/09/831,047C
CURRENT FILING DATE: 2001-05-03
PRIOR PLICATION NUMBER: 60/107,478
PRIOR PLICATION NUMBER: 60/107,478
PRIOR FILING DATE: 1998-11-06
NUMBER OF SEQ ID NOS: 121
SOUTHWER PATENTING NOS: 121
SOUTHWER PATENTING NOS: 121
                                                                                                             Gaps
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                                                Score 20; DB 22; Length 5;
Pred. No. 4.2e+06;
0: Mismatches 2; Indels
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Pred. No. 4.2e+06;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TYPE: PRT
; ORGANISM: peptido-mimetic of a Lewis antigen
US-09-831-047C-114
                                                                                                        0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-894-594-11
; Sequence 11, Application US/09894594
; GENERAL INFORMATION:
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OTHER INFORMATION: Novel Sequence
                                                  87.0%;
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                                                     Query Match
Best Local Similarity 60.0
Matches 3; Conservative
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Best Local Similarity
Matches 3; Conserv
US-09-831-047B-114
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US-09-831-047C-114
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LENGTH: 5
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CURRENT FILING DATE: 2000-01-19
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Matches 5; Conservative
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BRIAN, WEST
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APPLICANT:
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                                                                                                                                                    SEQ ID NO 1
LENGTH: 5
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APPLICANT: Baxter, John D.
APPLICANT: Baxter, John D.
APPLICANT: Darimont, Beatrice
APPLICANT: Darimont, Beatrice
APPLICANT: Pletterick, Robert J.
APPLICANT: Rushner, Peter J.
APPLICANT: Rushner, Peter J.
APPLICANT: Wagner, Richard L.
APPLICANT: Wagner, Richard L.
APPLICANT: Wagner, Reith R.
TITLE OF INVENTION: METHODS AND COMPOUNDS FOR MODULATING NUCLEAR RECEPTOR
TITLE OF INVENTION: COACTIVATOR BINDING
FILE REFERENCE: UCAL-253/02US
CURRENT APPLICATION NUMBER: US/09/281,717
CURRENT PILING DATE: 1999-03-30
EARLIER PILING DATE: 1999-03-30
NUMBER OF SEQ ID NOS: 51
SOFTWARE: PatentIn Ver. 2.1
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TITLE OF INVENTION: MATERIALS AND METHODS RELATING TO
TITLE OF INVENTION: INHIBITING THE INTERACTION OF p53 AND MDM2
FILE REFERENCE: MEMB25.001APC
CURRENT APPLICATION NUMBER: US/09/403,440A
                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence:peptide OTHER INFORMATION: Where Xaa may be any amino acid
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CURRENT APPLICATION NUMBER: US/09/214,371B CURRENT FILING DATE: 1999-03-26 PRIOR APPLICATION NUMBER: PCT/EP97/03549 PRIOR FILING DATE: 1997-07-04 NUMBER OF SEQ ID NOS: 83 SOFTWARE: Patentin Ver. 2.0
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LOCATION: (2)..(3)
OTHER INFORMATION: Xaa = Any Amino Acid
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nes 5; Conserv
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                                                                                                            SOFTWARE: 1
SEQ ID NO 2
LENGTH: 5
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Matches
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APPLICANT: YAMMOTO, KEITH
APPLICANT: GEISTLINGER, TIMOTHY
APPLICANT: GEISTLINGER, TIMOTHY
APPLICANT: GEISTLINGER, TIMOTHY
APPLICANT: ARNOLD, JAMES
APPLICANT: RUNTA, IRWIN
TITLE OF INVENTION: METHODS AND COMPOUNDS FOR MODULATING NUCLEAR RECEPTOR ACTIVITY
FILE REFERENCE: 9811-016-999
FULE REFERENCE: 9811-016-999
FULE REPERBATION NUMBER: US 60/079,965
PRIOR APPLICATION NUMBER: US 60/079,965
PRIOR APPLICATION NUMBER: US 60/113,146
PRIOR FILING DATE: 1998-03-30
PRIOR FILING DATE: 1999-03-30
PRIOR FILING DATE: 1999-03-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 82.6%; Score 19; DB 18; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 5; Conservative 0; Mismatches 0; Indels
PRIOR APPLICATION NUMBER: PCT/GB98/01144
PRIOR FILING DATE: 1998-04-20
PRIOR APPLICATION NUMBER: GB 9708092.3
PRIOR PILING DATE: 1997-04-22
NUMBER OF SEQ ID NOS: 12
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Xaa = any amino acid US-09-403-440A-1
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GENERAL INFORMATION:
APPLICANT: GUY, RODNEY
APPLICANT: BAXTER, JOHN
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PETER, KUSHNER
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Matches 2; Conservative
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ORGANISM: Homo Sapien
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PCT-US02-23257-10
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Sequence 791, Application PC/TUS0103988
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION OF MEMBRANE
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
CURRENT APPLICATION NUMBER: PCT/US01/03988
CURRENT APPLICATION NUMBER: 09/515,965
PRIOR APPLICATION NUMBER: 09/515,965
PRIOR APPLICATION NUMBER: 09/515,965
PRIOR APPLICATION NUMBER: 2000-02-29
SOFTWARE FARENES FARENES FOR WINDOWS VERSION 3.0
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                                                                APPLICANT: Roggen, Erwin
APPLICANT: Roggen, Erwin
APPLICANT: Svendsen, Allan
APPLICANT: Svendsen, Allan
APPLICANT: Svendsen, Allan
APPLICANT: Osten, Claus
TITLE OF INVENTION: PROTEIN VARIANTS HAVING MODIFIED IMMUNOGENICITY
FILE REFRENCE: 10021.204-US
CURRENT APPLICATION NUMBER: US/09/957,806A
CURRENT FILING DATE: 2001-09-21
NUMBER OF SEQ ID NOS: 248
SOFTWARE: Patentin version 3.1
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82.6%; Score 19; DB 23; Length 5;
Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2; Indels
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APPLICANT: Koister, Hubert
APPLICANT: Siddidi, Suhaib
APPLICANT: Little, Daniel
TITLE OF INVENTION: Capture Compounds, Collections Thereof
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65.2%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
US-09-957-806A-226
; Sequence 226, Application US/09957806A
; GENERAL INFORMATION:
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LENGTH: 5
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TITLE OF INVENTION: And Methods For Analyzing The Proteome And Complex TITLE OF INVENTION: Compositions
FILE REPRESENCE: 24743-2305
CURRENT APPLICATION NUMBER: PCT/US02/22821
CURRENT PILING DATE: 2002-07-16
PRIOR PELING DATE: 2001-07-16
PRIOR APPLICATION NUMBER: 60/306,019
PRIOR PELING DATE: 2001-08-21
PRIOR PELING DATE: 2001-08-21
PRIOR PELING DATE: 2001-08-21
PRIOR PELING DATE: 2002-03-11
NUMBER OF SEQ ID NOS: 149
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65.2%; Score 15; DB 1; Length 5;

Best Local Similarity 100.0%; Pred. No. 4.2e+06;

Matches 2; Conservative 0; Mismatches 0; Indels
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GENERAL INFORMATION:
APPLICANT: The Brigham and Women's Hospital, Inc.
TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: RECEPTOR LIGANDS
FILE REPRENCE: 50073/010W02
CURRENT APPLICATION NUMBER: PCT/US02/23257
CURRENT FILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: US 60/307,228
MINDER OF SETING DATE: 2001-07-23
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GENERAL INFORMATION:
APPLICANT: The Brigham and Women's Hospital, Inc.
TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: RECEPTOR LIGANDS
FILE REFERENCE: 50073/010W02
CURRENT APPLICATION NUMBER: PCT/US02/23257
CURRENT FILING DATE: 2002-07-23
PRIOR APPLICATION NUMBER: US 60/307,228
PRIOR FILING DATE: 2001-07-23
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SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 8
LENGTH: 5
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SOFTWARE: FastSEQ for Windows Version 4.0
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2, Conservative 0; Mismatches 0; Indels
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65.2%; Score 15; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
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TITLE OF INVENTION: ANALGESIC METHODS USING ENDOTHELIN
TITLE OF INVENTION: RECEPTOR LIGANDS
FILE REFERENCE: 50073/010W02
CURRENT APPLICATION NUMBER: PCT/US02/23257
CURRENT APPLICATION NUMBER: US 60/307,228
PRIOR FILING DATE: 2001-07-23
PRIOR FILING DATE: 2001-07-23
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
GENERAL INFORMATION:
ADPLICANT: The Brigham and Women's Hospital, Inc.
TITLE OF INVENTION: ANALGESIC WETHODS USING ENDOTHELIN
TITLE OF INVENTION: RECEPTOR LIGANDS
FILE REPREBENCE: 50073/0.00002
CURRENT APPLICATION NUMBER: PCT/US02/23257
CURRENT FILING DATE: 2002-07-23
FRIOR PELICATION UNMBER: US 60/307,228
FRIOR FILING DATE: 2001-07-23
NUMBER OF SEQ ID NOS: 14
SOOTWARE FASTESO FOR WINDOWS Version 4.0
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
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GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Artificial Sequence
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COTHER INFORMATION: Synthetic PCT-US02-23257-12
                                                                ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Synthetic PCT-US02-23257-11
                                                                                      FEATURE:
CTHER INFORMATION: Synthetic
PCT-US02-23257-10
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PCT-US02-23257-11
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; SEQ ID NO 10
                       LENGTH: 5
                                           TYPE: PRT
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Sequence 1, Application PC/TUS9815227
GENERAL INFORMATION:
APPLICANT: The New York Blood Center, Inc.
TITLE OF INVENTION: MONOSPECIFIC ANTIBODY REACTIVE WITH
TITLE OF INVENTION: MATRIX METALLOPROTEINASE CLEAVAGE PRODUCTS OF FIBRIN(OGEN)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ō
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                                                                                                                                  Sequence 13, Application PC/TUS0223257
Sequence 13, Application PC/TUS0223257
GENERAL INFORMATION:
APPLICANT: The Brigham and Women's Hospital, Inc.
TITLE OF INVENTION: RECEPTOR LIGANDS USING ENDOTHELIN
TITLE OF INVENTION: RECEPTOR LIGANDS
FILE REPERBNES: 50073/010002
CURRENT PAPLICATION NUMBER: 2002-07-23
PRIOR PILING DATE: 2002-07-23
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FRASESQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 65.2%; Score 15; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Ile at position 3 is alle PCT-USO2-23257-13
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APPLICATION NUMBER: PCT/US98/15227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Hoffmann & Baron, LLP STREET: 350 Jericho Turnpike
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/900,895
FILING DATE: 7/25/97
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SYSTEM: MS-DOS
WordPerfect
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: Synthetic
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
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                                                                                                                         PCT-US02-23257-13
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LENGTH: 5
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4 LW 5
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TELECOMMUNICATION INFORMATION:

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GENERAL INFORMATION:
APPLICANT: TRIMERIS, INC.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: PROPERTIES
FILE REFERENCE: 7872-052-228
CURRENT APPLICATION NUMBER: PCT/US99/11219
CURRENT PILING DATE: 1999-05-20
NUMBER OF SEQ ID NOS: 1542
SOFTWARE: FastSEQ for Windows Version 3.0
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COUNTRY: USA
ZIP: 20005

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/585,141
TITM DATE: 19901009
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GENERAL INFORMATION:
APPLICANT: Schramm, Wolfgang
APPLICANT: Schramm, Hans J.
TITLE OF INVENTION: Agent For Inhibiting Symmetrical
TITLE OF INVENTION: Proteins, In Particular Enzymes
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Venable, Baetjer, Howard & Civiletti
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1201 New York Avenue N.W., Suite 1000
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65.2%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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COTHER INFORMATION: Core polypeptide
PCT-US99-11219-791
TELEPHONE: (516) 822-3550
TELEPAX: (516) 822-3582
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                           ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
PCT-US98-15227-1
                                                                                                                 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREET: 1201 New CITY: Washington STATE: DC
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LENGTH: 5
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US-07-608-569-17
Sequence 17, Application US/07608569
GENERAL INFORMATION:
APPLICANT: Tainer, John A
APPLICANT: Tainer, John A
APPLICANT: Ruhn, Leslie
APPLICANT: Boissinct, Maurice
APPLICANT: Brisher, Cindy
APPLICANT: Griffin, John H
APPLICANT: Mallembach, Guy T
APPLICANT: Mal
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/608,569
FLING DATE: 19901102
CLASSIFICATION 1424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 05/07/608,569
FLING DATE: 01-NOV-1990
ATTONNEY/AGBNT IRFORMATION:
NAME: Fitting Thomas
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: 34,163
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEBRANCE 619-554-6312
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
PRIOR APPLICATION DATA:
APPLICATION NUMBER: WO PCT/EP90/00219
FILING DATE: 09-FEB-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: DE P 39 04 040.2
FILING DATE: 10-FEB-1989
ATTORNEY/AGENT INFORMATION:
NAME: Ihnen, Jeffrey L.
REGISTRATION NUMBER: 28957
REGISTRATION NUMBER: 18644-96040
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAX: 202-962-8100
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 INFORMATION FOR SEQ ID NO: 11: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 5 amino acids TYPE: AMINO ACID
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; HYPOTHETICAL: YES
US-07-585-141-11
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US-08-344-185B-454
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APPLICANT: SIMONS, TO SIMONS, T.J.
APPLICANT: BULOT-MATIAS, T.J.
APPLICANT: DAWSON, GEORGE G.
APPLICANT: SCHAMUDER, GEORGE G.
APPLICANT: SCHAMUDER, GEORGE G.
APPLICANT: LEARY, THOMAS P.
APPLICANT: LEARY, THOMAS P.
APPLICANT: MUSRAHOFF, ANTHONY T.
APPLICANT: MUSRAHWAR ISA
TITLE OF INVENTION: NON-A, NON-B, NON-C, NON-C, NON-E
TITLE OF INVENTION: HEPATITIS REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 539
CORRESPONDENCE ADDRESS:
STREET: ONE HUNDRED ABBOTT PARK ROAD
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                                                                                                                                                                                             65.2%; Score 15; DB 3; Length 5; ilarity 100.0%; Pred. No. 4.2e+06; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/344,185B
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 100, Application US/08344185B GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY AGENT INFORMATION:
NAME: POREMESKI, PRISCILLA E.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-97-6365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: 708-937-6365
TELEPAX: 708-938-2623
INPORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Floppy disk
                       SEQUENCE CHARACTERISTICS:
  INFORMATION FOR SEQ ID NO:
                                       LENGTH: 5 amino acids
TYPE: AMINO ACID
TOPOLOGY: linear
                                                                                                         MOLECULE TYPE: peptide FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
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COMPUTER READABLE FORM:
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CLASSIFICATION: 435
                                                                                                                                                                                                                    Best Local Similarity
Matches 2; Conserv
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RESULT 25

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GENERAL INFORMATION:
GENERAL INFORMATIONS, JOHN N.
APPLICANT: SIMONS, JOHN N.
APPLICANT: DAWSON, GENERE J.
APPLICANT: DAWSON, GENERE J.
APPLICANT: DESAI, SURESH M.
APPLICANT: DESAI, SURESH M.
APPLICANT: LERRY, THOMAS P.
APPLICANT: BUJJK, S.L.
APPLICANT: MUSHAPHWAR, ISA K.
TITLE OF INVENTION: NON-A, NON-B, NON-C, NON-D, NON-E
TITLE OF INVENTION: HEPATITIS REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 539
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                           APPLICANT: LEARY, THOMAS P.
APPLICANT: MUERHOFF, ANTHONY T.
APPLICANT: BUILK, S.L.
APPLICANT: MUSHAHWAR, ISA K.
TITLE OF INVENTION: NON-A, NON-B, NON-C, NON-D, NON-E
TITLE OF INVENTION: HEPATITIS REAGENTS AND METHODS FOR THEIR USE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 65.2%; Score 15; DB 7; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 539
CORRESPONDENCE ADDRESS:
ADDRESSE: ABBOTT LABORATORIES
STREET: ONE HUNDRED ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: ABBOTT LABORATORIES
STREET: ONE HUNDRED ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/344,185B
; Sequence 454, Application US/08344185B ; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Application US/08344185C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PC-DOS/MS-DOS
                                                    SIMONS, JOHN N.
PILOT-MATIAS, T.J.
DAWSON, GEORGE J.
SCHLAUDER, GEORGE G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: POREMISKI, PRISCILLA E.
TELECOMMINICATION INFORMATION:
TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-
                                                                                                                                                           DESAI, SURESH M.
LEARY, THOMAS P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         708-938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-344-1858-454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION: 435
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ZIP: 60064-3500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               USA
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US-08-344-185C-100
                                                         APPLICANT:
                                                                                                                                   APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 LW 5
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LENGTH: 5 amino acids
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APPLICANT: BLOT-MATIAS, T.J.
APPLICANT: BLOT-MATIAS, T.J.
APPLICANT: SCHLAUDER, GEORGE J.
APPLICANT: SCHLAUDER, GEORGE G.
APPLICANT: LEARY, THOMAS P.
APPLICANT: LEARY, THOMAS P.
APPLICANT: BUICK, S.L.
APPLICANT: WUSENOFF, ANTHONY T.
APPLICANT: MUSENATION: HEPATITIS REAGENTS AND METHODS FOR THEIR USE CORRESPONDENCE ADDRESSES:
ADDRESSEE: ABBOTT LABORATORIES
STREET: ONE HUNDRED ABBOTT PARK ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 65.2%; Score 15; DB 7; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
                                              ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/344,185C
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ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/344,185C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 454, Application US/08344185C GENERAL INFORMATION:
                                                                                                                                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
TELEFAX: 708-938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: POREMBEKI, PRISCILLA E.
TELECOMMINICATION INFORMATION:
TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 708-938-2623
INPORMATION FOR SEQ ID NO: 100:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     454:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: protein
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3Y: linear
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USA
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                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
NUMBER OF SEQUENCES: 116
CORRESPONDENCE ADDRESS:
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                                                                                              Query Match 65.2%; Score 15; DB 7; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
COUNTRY: USA
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 65.2%; Score 15; DB 8; Le
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SSEE: ABBOTT LABORATORIES D377/AP6D
1: 100 ABBOTT PARK ROAD
ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      5527.PC.01
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/424,550A
                                                                                                                                                                                                                                                                                                                                                               JOHN N. SIMONS
TAMI J. PILOT-MATIAS
GEORGE J. DAMSON
GEORGE G. SCHLAUDER
SURESH M. DESAI
THOMAS P. LEARY
ANTHONY SCOTT MUERHOFF
JAMES C. ERKER
SHERI L. BULJK
                                                                                                                                                                                                                                                                                  RESULT 28
US-08-424-550A-228
; Sequence 228, Application US/08424550A
; GENERAL INFORMATION:
; APPLICANT: JOHN N. SIMONS
; APPLICANT: TAMI J. PILOT-MATIAS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33, 207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: 55:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
TELEPAX: 708-938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 228:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-344-185C-454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-424-550A-228
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US-08-424-550B-228
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                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 716
CORRESPONDENCE ADDRESS:
ADDRESSE: ABBOTT LABORATORIES D377/AP6D
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             .
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GENERAL INFORMATION:
APPLICANT: Kishore, Ganesh M.
APPLICANT: Arondel, Vincent J.
APPLICANT: Arondel, Vincent J.
APPLICANT: Gomerville, Christopher R.
APPLICANT: Gomerville, Christopher R.
TITLE OF INVENTION: Altered Linolenic and Linoleic Acid
TITLE OF INVENTION: Content in Plants
NUMBER OF SEQUENCES: 72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 E: Lawrence M. Lavin, Jr., Monsanto Co., BB4F 700 Chesterfield Parkway North
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Ouery Match
65.2%; Score 15; DB 8; Length 5;
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE:
CLASSIFICATION: 435435
ATTONREY, AGENT INFORMATION:
NAME: PORENBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION:
TELEPHONE: 708-939-6365
TELEFAX: 708-938-2623
                                   JOHN N. SIMONS
TAMI J. PILOT-MATIAS
GEORGE J. DAWGON
GEORGE G. SCHLAUDER
SURESH M. DESAN
THOMAS P. LEARY
ANTHONY SCOTT MUERHOFP
JAMES C. ERKER
SHERI L. BUIJK
ISA K. MUSHAHWAR
Application US/08424550B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 228:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    : 5 amino acids
amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Lawrence N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: St. Louis
STATE: Missouri
                                                                                                                                                                                                                                                                                                                                                                                                                                             60064-3500
                        GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 30
US-08-432-147-47
                                                                                APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                               APPLICANT:
APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
  ; Sequence 228,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET:
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                                                                 APPLICANT:
                                                                                                                                                                                                              APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match

65.2%; Score 15; DB 8; Length 5;
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
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OPERATING SYSTEM: PC_DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                D377/AP6D
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
OPENATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/432,147
FILING DATE: 05-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
RAPPLICATION NUMBER: US/08/467,344A
FILING DATE: 07-Jun-1995
CLASSIFICATION: <URKnown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Lavin Jr., Lawrence M. REGISTRATION NUMBER: 30,768
REFERENCE/DOCKET NUMBER: 38-21(10650)A TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: ABOUT LABORATORIES
STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
GENERAL INFORMATION:
HAPLICANT: JOHN N. SIMONS
GEORGE J. DAWSON
GEORGE J. DAWSON
GEORGE J. DAWSON
HAPPING STRESH M. DESAI
THOMAS P. LEARY
ANTHONY SCOTT MUERHOFF
JAMES C. ERKER
SHERI L. BUIJK
HAPPING STREER
SHERI L. BUIJK
HAPPING STREER
HAPP
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FILING DATE: 04-FEB-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                               FILING DATE: 05-MAY-1995
CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCTUS94/01321
FILING DATE: 04-FEB-1994
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (314)537-6047
INFORMATION FOR SEQ ID NO: 47:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (314)537-6670
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 60064-3500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; MOLECULE TYPE: peptide US-08-432-147-47
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4 LW 5
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Query Match 65.2%; Score 15; DB 8; Le
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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STREET: 100 ABBOTT PARK ROAD
CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                          RESULT 33
US-08-488-445A-228
i Sequence 228, Application US/08488445A
i GENERAL INFORMATION:
APPLICANT: JOHN N. SIMONS
APPLICANT: THAM J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: THOMAS P. LEARY
APPLICANT: THOMAS P. LEARY
APPLICANT: ANTHONY SCOTT MUERHOFF
APPLICANT: SHERI L. BUIJK
APPLICANT: TON V. MISHAHWAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO: 228:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                     2; Conservative
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                                ; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-486-749-228
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TOPOLOGY: linear
          amino acid
                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                          Matches
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APPLICANT: JOHN N. SIMONS
APPLICANT: GEORGE J. DAMSON
APPLICANT: GEORGE J. DAMSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: STRESH M. DESAI
APPLICANT: ANTHONAS P. LEARY
APPLICANT: ANTHONY SCOTT MUERHOFF
APPLICANT: JAMES C. EKER
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: BHR PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIJ Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,749
                                           ATTORNEY/AGENT INFORMATION:
NAME: POREMEST, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-938-5623
INFORMATION FOR SEQ ID NO: 228:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
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                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein

SEQUENCE DESCRIPTION: SEQ ID NO: 228:
US-08-467-344A-228
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/424,550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 5527.PC.01
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 228, Application US/08486749 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: POREMBSKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/POCKET NUMBER: 5527.
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 228: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 32
US-08-486-749-228
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                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEFATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
NUMBER OF SEQUENCES: 116
CORRESPONDENCE ADDRESS:
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65.2%; Score 15; DB 8; Length 5; 100.0%; Pred. No. 4.2e+06; vative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: USA
ZIP: 60064-3500
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,445A
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US-08-488-446-228
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COUNTRY:
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GENERAL INFORMATION:
APPLICANT: HIGGER, CHRISTOPH-STEPH
APPLICANT: DINKELBORG, LUDGER
APPLICANT: KRAMP, WOLGANG
APPLICANT: KRAMP, WOLGANG
APPLICANT: SCHIEK, HANS-MARTIN
TITLE OF INVENTION: CHELATING AGENTS OF THE TYPE XNISIOI FOR RADIOACTIVE I
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
ADDRESSEE: FLIESTER, DUBB, MEYER & LOVEJOY
STREET: FOUR EMBASCAGERO Center, Suite 400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ;
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REAGENTS AND METHODS FOR THEIR USE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC_Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/488,446
                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: ABBOTT LABORATORIES D377/AP6D STREET: 100 ABBOTT PARK ROAD CITY: ABBOTT PARK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
NAME: POREMESKI, PRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION INFORMATION:
TELEFAX: 708-938-2635
INFORMATION FOR SEQ ID NO: 228:
SEQUENCE CHARACTERISTICS:
                                    APPLICANT: JOHN N. SIMONS
APPLICANT: JOHN N. SIMONS
APPLICANT: TANI J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: SURESH M. DESAI
APPLICANT: THOMAS P. LEARY
APPLICANT: ANTHONY SCOTT MUERHOFF
APPLICANT: JAMES C. ERKER
APPLICANT: SHERI L. BUIJK
APPLICANT: SHERI L. BUIJK
APPLICANT: ISA K. MUSHAHWAR
TITLE OF INVENTION: NON-A, NON-B. NC
TITLE OF INVENTION: REAGENTS AND MET
NUMBER OF SEQUENCES: 716
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550
Application US/08488446
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: amino acid
; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-488-446-228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: Four Embarce
CITY: San Francisco
STATE: CA
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Best Local Similarity 1
'-hem 2; Conservat
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  ; Sequence 228, Applica ; GENERAL INFORMATION:
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US-08-530-325-14
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RESULT 36
US-08-530-127-14
Sequence 14, Application US/08530327
Sequence 14, Application US/08530327
Sequence 14, Application US/08530327
Sequence 14, Application US/08530327
Sequence 14, Application
APPLICANT: BINKELEDROG, LUDGER
APPLICANT: SCHULTZE, PAUL-EBERHARD
APPLICANT: NOLL, BERNHARD
APPLICANT: NOLL, BERNHARD
APPLICANT: SCHULTZE, PAUL-EBERHARD
APPLICANT: SCHULTZE, PAUL-EBERHARD
APPLICANT: NOLL, BERNHARD
CORRESPONDENCES: 16
NUMBER OF SEQUENCES: 16
CORRESPONDENCES: 6
ADDRESSES: FLIESLER, DUBB, MEYER & LOVEJOY
STREET: FOUR Embarcadero Center, Suite 400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                               REFERENCE/DOCKET NUMBER: WBLT 2005 (WO) PA
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-362-3800
                                                                                                                                                                            CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DE94/00371
FILING DATE: 29-MAR-1994
APPLICATION NUMBER: P 43 11 023.1
FILING DATE: 31-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Dubb, Hubert E
REGISTRATION NUMBER: 25,329
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUFTWARE: LESTEM: DOS
SUFTWARE: FastSEG Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/530,327
FILING DATE:
CLASSIFICATION: 424
                                                                     OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ VERSION 1.5
SOFTWART APPLICATION DATA:
APPLICATION NUMBER: US/08/530,325
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COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                             IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 415-362-2928
                            Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE: peptide HYPOTHETICAL: NO
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                   OPERATING SYSTEM:
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US-08-530-325-14
                                                  COMPUTER:
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TYPE: PRT
ORGANISM: Artificial Sequence
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        GENERAL INFORMATION:
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APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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65.2%; Score 15; DB 13; Length 5;
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           65.2%; Score 15; DB 9; Length 5; 100.0%; Pred. No. 4.2e+06; vative 0; Mismatches 0; Indels
                                                                                                                                                                WBLT 2007 (WO) PA
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CURRENT APPLICATION NUMBER: US/08/981,122A
CURRENT FILING DATE: 1997-12-18
EARLIER APPLICATION NUMBER: UP 7-176904
EARLIER APPLICATION NUMBER: PCT/JP96/01734
EARLIER FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: PATENTIN Ver. 2.0
                                   FILING DATE: 29-MRR-1994
APPLICATION WUMBER: P 43 11 021.5
FILING DATE: 31-MRR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Dubb, Hubert E
REGISTRATION WUMBER: 25,329
REFERENCE/DOCKET NUMBER: WBLT 2007 (
                      PCT/DE94/00369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 37
US-08-981-122-15
; Sequence 15, Application US/08981122A
; GENERAL INFORMATION:
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US-08-981-122-38
; Sequence 38, Application US/08981122A
                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-362-3800
                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2; Conservative
                                                                                                                                                                                                                        TELEFAX: 415-362-2928
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MOLECULE TYPE: peptide
HYPOTHETICAL: NO
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 2; Conserv
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LENGTH: 5
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APPLICANT: Attendary Constitution of APPLICANT: Attendary APPLICANT: Attendary APPLICANT: Attendary APPLICANT: Attendary APPLICANT: Attendary APPLICANT: Attendary APPLICANT: AP
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APPLICANT: Barney, S.
APPLICANT: Guthrie, K.
APPLICANT: Menutka, G.
APPLICANT: Anner, M.
APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC PROPERTIES
GENERAL INFORMATION:
APPLICANT: Genencor International, Inc.
TITLE OF INVENTION: Novel EGIII-Like Enzymes, DNA Encoding
TITLE OF INVENTION: Novel EGIII-Like Enzymes, DNA Encoding
TITLE OF INVENTION: Such Enzymes and Methods for Producing Such Enzymes
FILE REFERENCE: GCS16-2-PCT
CURRENT APPLICATION NUMBER: US/09/284,327
CURRENT PILING DATE: 1999-04-10
NUMBER OF SEQ ID NOS: 41
SOFTWARE: FASELSEQ for Windows Version 3.0
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GENERAL INFORMATION:
TITLE OF INVENTION: Novel Prokaryotic Polynucleotides,
TITLE OF INVENTION: Polypeptides and Their Uses
FILE REFERENCE: P50596
CURRENT APPLICATION NUMBER: US/09/308,823A
CURRENT FILING DATE: 1999-05-25
NUMBER OF SEQ ID NOS: 661
SOFTWARE: FRAESEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 65.2%; Score 15; DB 16; L
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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CURRENT APPLICATION NUMBER: US/09/350,641

CURRENT FILING DATE: 1999-07-09

PRIOR APPLICATION NUMBER: 09/315,304

PRIOR FILING DATE: 1999-05-20

PRIOR FILING DATE: 1998-05-20

PRIOR FILING DATE: 1998-05-20

NUMBER OF SEQ ID NOS: 1589

SOFTWARE: FASESEQ FOr Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Staphylococcus aureus
                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Synthetic
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Best Local Similarity
Thes 2; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 43
US-09-308-823A-582
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US-09-350-641-791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid OTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ·;
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TITLE OF INVENTION: The Use of Consensus Sequences for Targeted Homologous
TITLE OF INVENTION: The Use of Consensus Sequences for Targeted Homologous
TITLE OF INVENTION: Gene Isolation and Recombination in Gene Families
FILE REFRENCE: A-65678-1/FFT/NBC
CURRENT APPLICATION NUMBER: US/09/209,676
CURRENT FILING DATE: 2002-06-18
PRIOR APPLICATION NUMBER: US 60/070,734
PRIOR FILING DATE: 1997-12-11
NUMBER OF SEQ ID NOS: 95
SOFTWARE: PatentIn Ver. 2.0
                                                                                                     APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
IITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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US-09-209-676-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        65.2%; Score 15; DB 13; Length 5; ilarity 100.0%; Pred. No. 4.2e+06; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                              FILE REPERENCE:
CURRENT APPLICATION NUMBER: US/08/981,122A
CURRENT FILING DATE: 1997-12-18
EARLIER APPLICATION NUMBER: UP 7-176904
EARLIER FILING DATE: 1995-06-21
EARLIER FILING DATE: 1995-06-21
EARLIER FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
                                                     Sequence 51, Application US/08981122A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 30, Application US/09209676; GENERAL INFORMATION:
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Matches 2; Conserva
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                                     US-08-981-122-51
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FEATURE:
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US-09-410-568-9
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      SEQ ID NO 791
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LENGTH: 5
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APPLICANT: Anwer, M.
APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: 1972-067
CURRENT PEDIAGRATION NUMBER: US/09/350,641B
PRIOR PLING DATE: 1999-07-09
PRIOR PLING DATE: 1999-05-20
PRIOR PLING DATE: 1999-05-20
PRIOR PLING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1757
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 791
LENGTH: S
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APPLICANT: Merutka, G.
APPLICANT: Anwer, M.
APPLICANT: Anwer, M.
APPLICANT: Anwer, M.
TITLE OF INVENTION: PROPERTIES
TITLE OF INVENTION: PROPERTIES
TILL OF INVENTION: PROPERTIES
TILL OF TINGENT APPLICATION NUMBER: US/09/350,641C
CURRENT APPLICATION NUMBER: 09/315,304
PRIOR PRILIAG DATE: 1999-05-20
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1999-05-20
PRIOR FILING DATE: 1998-05-20
PRIOR FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1757
SOFTWARE: PASISEQ for Windows Version 3.0
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; APPLICANT: Barney, S.
                                                                                   ; OTHER INFORMATION: Core polypeptide US-09-350-641-791
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; OTHER INFORMATION: Core polypeptide
US-09-350-641B-791
                    TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
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ORGANISM: Artificial Sequence
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APPLICANT: Bhatia, Ajay
APPLICANT: Bhatia, Ajay
APPLICANT: Shatia, Ajay
APPLICANT: Stomberg, Hang
APPLICANT: Fang, Hang
APPLICANT: Stromberg, Erica Jean
APPLICANT: Stromberg, Erica Jean
APPLICANT: Stromberg, Brica Jean
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
CURRENT APPLICATION NUMBER: US/09/410,568
CURRENT FILING DATE: 1999-10-01
NUMBER OF SEQ ID NOS: 109
SOFTWARE FASTSEQ FOR Windows Version 3.0
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APPLICANT: Bhatia, Ajay
APPLICANT: Bhatia, Ajay
APPLICANT: Skeiky, Yasir
APPLICANT: Fling, Steve
APPLICANT: Fang, Hang
APPLICANT: From Shyins
APPLICANT: Stromberg, Erica Jean
APPLICANT: Enghart, Susan E.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR TREATMENT AND
FILE REFERENCE: 469C3
                                                                                                                                                                     65.2%; Score 15; DB 17; Length 5; 100.0%; Pred. No. 4.2e+06; tive 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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CURRENT FILING DATE: 1999-10-22
NUMBER OF SEQ ID NOS: 168
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                   FEATURE:
, OTHER INFORMATION: Core polypeptide
US-09-350-641C-791
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US-09-426-571-9
: Sequence 9, Application US/09426571
: GENERAL INFORMATION:
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ORGANISM: Chlamydia trachomatis
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LENGTH: 5
TYPE: PRT
ORGANISM: Artificial Sequence
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Best Local Similarity 100..
Thes 2; Conservative
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1 LW 2
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APPLICANT: Skeiky, Yasir
APPLICANT: Fling, Stave of APPLICANT: Maisonneuve, Jeff
ITTLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
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                                   Query Match 65.2%; Score 15; DB 18; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
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CURRENT APPLICATION NUMBER: US/09/454,684A
CURRENT FILING DATE: 1999-12-03
NUMBER OF SEQ ID NOS: 310
SOFTWARE: FastSEQ for Windows Version 3.0/4.0
SEQ ID NO 9
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILE REFERENCE: 210121.469C4
CURRENT APPLICATION WUMBER: US/09/454,684
CURRENT FILING DATE: 1999-12-03
NUMBER OF SEQ ID NOS: 303
SOFTWARE: FastSEQ for Windows Version 3.0/4.0
SEQ ID NO 9
LENOTH: 5
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                                                                                                                                                                                                                                                                                    Sequence 9, Application US/09454684; GENERAL INFORMATION:
APPLICANT: Probst, Peter
APPLICANT: Bhatia, Ajay
APPLICANT: Skelky, Yasair
APPLICANT: Fling, Steve;
APPLICANT: Maisonneuve, Jeff
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Best Local Similarity
Matches 2; Conserv
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US-09-454-684A-9
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US-09-454-684-9
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US-09-426-571-9
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APPLICANT: Antczak, J.
APPLICANT: Delmedico, M.
APPLICANT: Delmedico, M.
APPLICANT: Erickson, J.
APPLICANT: Exickson, J.
APPLICANT: Sista, P.
APPLICANT: Sista, D.
APPLICANT: Sis
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                                   Sequence 25, Application US/09457198A

GENERAL INFORMATION:
APPLICANT: Christine, Andrews A.
APPLICANT: Hartnett, James R.
ITLE OF INVENTION: Thermostable Enzyme for Nucleic Acid Detection
FILE REPERENCE: Thermostable/nucleic acid detection
CURRENT APPLICATION NUMBER: US/09/457,198A

CURRENT FILING DATE: 1999-12-06
FRIOR FILING DATE: 1999-07-21
NUMBER OF SEQ ID NOS: 28

SOFTWARE: Patentin Ver. 2.1

SEQ ID NO 25

LENGTH: 5
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65.2%; Score 15; DB 18; Length 5;
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
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CURRENT FILING DATE: 2000-02-29
FRICR APPLICATION WUMBER: 09/315,304
PRIOR APPLICATION WUMBER: 09/082,279
PRIOR PILING DATE: 1998-05-20
PRIOR FILING DATE: 1998-05-20
SUMMBER OF SEQ ID NOS: 1589
SOFTWARE: FREESEQ FOR WINDOWS Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-515-965A-791
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US-09-515-965-791
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US-09-457-198A-25
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LENGTH: 5
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Gaps

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APPLICANT: Kudryk, Bohdan

TITLE OF INVENTION: MONOSPECIFIC ANTIBODY REACTIVE WITH MATRIXMETALLOPROTEINASE CLEAV

TITLE OF INVENTION: PRODUCTS OF FIBRIN (OGEN)

FILE REPRESENCE: 454-16CTP

CURRENT APPLICATION NUMBER: US/09/536,998

CURRENT FILING DATE: 2001-04-25

PRIOR PILING DATE: 1997-07-25

NUMBER OF SEQ ID NOS: 3

SOFTWARE: PATENTIN VERSION 3.0

SEQ ID NO 1

LENGTH: 5
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US-09-588-419-9
; Sequence 9, Application US/09598419
; Sequence 9, Application US/09598419
; GENERAL INFORMATION:
; APPLICANT: Scholler, John
; TITLE OF INVENTION: COMPOUNDS AND METHODS FOR TREATMENT AND
; TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
; TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
; TITLE OF INVENTION: US/09/598,419
; CURRENT APPLICATION NUMBER: US/09/598,419
; CURRENT FILING DATE: 2000-06-20
; NUMBER OF SEQ ID NOS: 357
; SOFTWARE: FASISEQ for Windows Version 3.0/4.0
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APPLICANT: Ropp, Traci M.
APPLICANT: Swanson, Barbara A.
TITLE OF INVENTION: Novel Variant EGIII-Like Cellulase
TITLE OF INVENTION: Compositions
FILE REFERENCE: GC630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  65.2%; Score 15; DB 19; L
100.0%; Pred. No. 4.2e+06;
trive 0; Mismatches 0;
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100.0%; Pred. No. 4.2e+06;
ative 0; Mismatches 0;
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CURRENT FILING DATE: 2000-08-04
NUMBER OF SEQ ID NOS: 86
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 30
LENGTH: 5
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US-09-632-426-30
US-09-632-426-30
Sequence 30, Application US/09632426
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT; ORGANISM: Homo sapiens
US-09-536-998-1
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Best Local Similarity
2, Conserve
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Best Local Similarity
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GENERAL INFORMATION:
APPLICANT: Antczak, J.
APPLICANT: Delmedico, M.
APPLICANT: Existen, J.
APPLICANT: Existen, J.
APPLICANT: Sista, P.
APPLICANT: Sista, Sista, Sista, P.
APPLICANT: Sista, Sista, P.
APPLICANT: Sista, Sista, Sista, P.
APPLICANT: Sista, Sista, Sista, Sista, P.
APPLICANT: Sista, Sista, Sista, Sista, P.
APPLICANT: Sista, Sis
                                      APPLICANT: Sista, P.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR INHIBITION OF MEMBRANE FUSION-
TITLE OF INVENTION: ASSOCIATED EVENTS INCLUDING RSV TRANSMISSION
FILE REFERENCE: 7872-073
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100.0%; Pred. No. 4.2e+06;
ative 0; Mismatches 0;
                                                                                                                                                FILE NEFERENCE (0.75 CV) 3
CURRENT PELLOXION NUMBER: US/09/515,965A
CURRENT FILING DATE: 2000-02-29
PRIOR APPLICATION NUMBER: 09/315,304
PRIOR FILING DATE: 1999-05-20
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1994
SEQ ID NO 791
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Sequence 1, Application US/09536998
GENERAL INFORMATION:
APPLICANT: New York Blood Center
APPLICANT: Bini, Alessandra
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ORGANISM: Artificial Sequence
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         Lambert, D.
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Best Local Similarity
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Matches 2; Conserv
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         APPLICANT:
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APPLICANT: Mitchinson, Colin
APPLICANT: Phillips, Jay Ian
TITLE OF INVENTION: Novel Variant EGIII-Like Cellulase
TITLE OF INVENTION: Compositions
FILE REFERENCE: GC631
CURRENT APPLICATION NUMBER: US/09/632,570
CURRENT FILING DATE: 2000-08-04
NUMBER OF SEQ ID NOS: 64
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 5
                                                                                             DB 20; Le
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             65.2%; Score 15; DB 20; I 100.0%; Pred. No. 4.2e+06;
                                                                                         ilarity 100.0%; Pred. No. 4.2
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: WANG, Bryan S.
APPLICANT: WANG, Bryan S.
APPLICANT: PABO, CALL O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243A
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN Ver. 2.0
                LOCATION: (1)...(5)
CTHER INFORMATION: Xaa = Asn or Gln US-09-632-426-30
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OTHER INFORMATION: Xaa = Asn or Gln
OTHER INFORMATION: BOX1
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                                                                                                                                                                                                                                                                                                           Sequence 30, Application US/09632570
GENERAL INFORMATION:
APPLICANT: Gualfetti, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: PRT
ORGANISM: Artificial Sequence
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                                                                         Query Match
Best Local Similarity
NAME/KEY: VARIANT
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APPLICANT: Conjuchem, Inc.
APPLICANT: Bridon, Dominique
APPLICANT: Bridon, Dominique
APPLICANT: Bridon, Peter
APPLICANT: Milner, Peter
APPLICANT: Holmes, Darren
APPLICANT: Thibaudeau, Karen
APPLICANT: THIBAUTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: COMPONENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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CURRENT APPLICATION NUMBER: US/09/657,276
CURRENT FILING DATE: 2000-09-07
PRIOR APPLICATION NUMBER: 60/134,406
PRIOR FILING DATE: 1999-05-17
PRIOR FILING DATE: 1999-09-10
PRIOR FILING DATE: 1999-09-10
PRIOR APPLICATION NUMBER: 60/159,783
PRIOR FILING DATE: 1999-10-18
NUMBER OF SEQ ID NOS: 1617
SEQ ID NO 723
                                                                                                                                                                                                                      APPLICANT: WANG, Bryan S.
APPLICANT: WANG, Bryan S.
TITLE OF INVENTION: DIMERIZING PEPTIDES
TITLE OF INVENTION: DIMERIZING M4-US2
CURRENT APPLICATION NUMBER: US/09/642,234
CURRENT FILING DATE: 2000-08-17
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR PILING DATE: 1999-08-11
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                                                                                                                                                 US-09-642-234-75; Sequence 75, Application US/09642234; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Artificial Sequence
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Best Local Similarity 100...
And 2; Conservative
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SOFTWARE: Patentin Ver. 2.0
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US-09-657-276-723
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4 LW 5
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LENGTH: 5
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65.2%; Score 15; DB 20; Length 5;

Query Match

Best Local Similarity

Query Match

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RESULT 64
US-09-657-276-1560
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                                 SEQ ID NO 734
LENGTH: 5
TYPE: PRT
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                  SOFTWARE:
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APPLICANT: Conjuchem, Inc.
APPLICANT: Bridon, Dominique
APPLICANT: Bridon, Dominique
APPLICANT: Bridon, Alan
APPLICANT: Holmes, Darren
APPLICANT: Holmes, Darren
APPLICANT: Holmes, Darren
APPLICANT: Thibaudeau, Karlon
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: COMPONENTS
TITLE OF INVENTION: COMPONENTS
FILE REFERENCE: 2110
CURRENT APPLICATION NUMBER: 08/09/07
PRIOR PILING DATE: 1999-09-10
PRIOR FILING DATE: 1999-09-10
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APPLICANT: Exin, Alan
APPLICANT: Exin, Alan
APPLICANT: Exin, Alan
APPLICANT: Exin, Alan
APPLICANT: Hibaudeau, Raren
APPLICANT: Hibaudeau, Raren
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: PROTECTION OF ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION: COMPONENTS
TITLE OF INVENTION: ENDOGENOUS THERAPEUTIC PEPTIDES FROM
TITLE OF INVENTION TOWBER: US/09/657,276
CURRENT APPLICATION NUMBER: 60/134,406
PRIOR APPLICATION NUMBER: 60/134,406
PRIOR APPLICATION NUMBER: 60/153,406
PRIOR PILING DATE: 1999-05-10
PRIOR PILING DATE: 1999-10-18
PRIOR PILING DATE: 1999-10-18
                                      Gaps
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                                      0; Indels
            Pred. No. 4.2e+06;
100.0%; Preu. ...
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ORGANISM: Artificial Sequence
                                      2; Conservative
          Best Local Similarity
Matches 2; Conserv
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US-09-657-276-734
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US-09-657-276-724
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APPLICANT: MORRELLS, HENRI
APPLICANT: MORRELLS, HENRI
APPLICANT: MARRELS, HENRI
TITLE OF INVENTION: BENZODIAZEPINES AND BENZOTHIAZEPINES DERIVATIVES AND
TITLE OF INVENTION: HESAG PEPTIDES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: AND USE
FILE REFERENCE: 2551-54
CURRENT APPLICATION NUMBER: US/09/762,270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: BILLOW: Alan
APPLICANT: BILLOW: Alan
APPLICANT: Milner, Peter
APPLICANT: Milner, Peter
APPLICANT: Milner, Peter
APPLICANT: Holmes, Darren
APPLICANT: Thibaudeau, Karen
TITLE OF INVENTION: PEPTIDASE ACTIVITY THROUGH CONJUGATION TO BLOOD
TITLE OF INVENTION: COMPONENTS
FILE REFERENCE: 2110
CURRENT APPLICATION NUMBER: US/09/657,276
CURRENT PILLOM DATE: 2000-09-07
PRIOR FILING DATE: 1999-09-10
PRIOR FILING DATE: 1999-09-10
PRIOR PRILICATION NUMBER: 60/153,406
PRIOR FILING DATE: 1999-09-10
PRIOR APPLICATION NUMBER: 60/159,783
PRIOR FILING DATE: 1999-09-10
PRIOR PILLING DATE: 1999-10-18
NUMBER OF SEQ ID NOS: 1617
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                                                                                                                                                                      ) OTHER INFORMATION: Description of Artificial Sequence: Synthetic; CTHER INFORMATION: Peptide US-09-657-276-734
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 1560, Application US/09657276; GENERAL INFORMATION:
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                                                                                                                 ORGANISM: Artificial Sequence
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Bridon, Dominique
Patentin Ver. 2.1
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APPLICANT: DEFLA, ERIK
APPLICANT: MAGREELS, HERRI
APPLICANT: MAGREELS, GEERT
APPLICANT: MAGREELS, GEERT
TITLE OF INVENTION: HEASG PEPTIDES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: AND USE
TITLE OF INVENTION: AND USE
TITLE OF INVENTION: AND USE
CURRENT APPLICATION NUMBER: US/09/762,270
CURRENT APPLICATION NUMBER: US/09/762,270
CURRENT PILING DATE: 1999-08-25
PRIOR APPLICATION NUMBER: EP 9897006231
PRIOR APPLICATION NUMBER: EP 98970062.9
PRIOR APPLICATION NUMBER: EP 98970062.9
PRIOR APPLICATION NUMBER: EP 99970062.9
PRIOR PILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 70
SOFTWARE PATEUT OF VERY PAT
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APPLICANT: MOEREELS, HENRI
APPLICANT: MAERTEELS, HENRI
APPLICANT: MAERTEENS, GEERL
TITLE OF INVENTION: BRACEDIAZEPINES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: AND USE
FILE REFERENCE: 2551-54
CURRENT APPLICATION NUMBER: US/09/762,270
CURRENT APPLICATION NUMBER: US/09-06
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PRIOR APPLICATION NUMBER: PCT/EP99/06231
PRIOR PLLICATION NUMBER: PCT/EP99/06231
PRIOR PLLICATION NUMBER: PP 98870186.8
PRIOR APPLICATION NUMBER: EP 98870186.8
PRIOR APPLICATION NUMBER: EP 99870062.9
PRIOR FILING DATE: 1999-03-29
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 70
SOFTWARE: PatentIn Ver. 2.1
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US-09-762-270-20
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APPLICANT: DEPLA, ERIK
APPLICANT: DEPLA, ERIK
APPLICANT: MORREELS, HENRI
APPLICANT: MARCHES, HENRI
TITLE OF INVENTION: BENZOIAZEPINES AND BENZOTHIAZEPINES DERIVATIVES AND
TITLE OF INVENTION: BENZOIAZEPINES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: AND USE
TITLE OF INVENTION: AND USE
TILLE OF INVENTION: AND USE
TILLE OF INVENTION: AND USE
TILLE OF INVENTION: DATE: 2011-02-06
CURRENY FILING DATE: 2011-02-06
PRIOR PLILING DATE: 1999-08-25
PRIOR FILING DATE: 1999-08-25
PRIOR FILING DATE: 1999-08-09-01
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 70
SOFTWARE: PARCHELIN VEY: 2.1
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GENERAL INFORMATION:
APPLICANT: DEPLA, ENER I
APPLICANT: MAERELS, HENRI
APPLICANT: MAERELS, HENRI
TITLE OF INVENTION: BENZODIAZEPINES AND BENZOTHIAZEPINES DERIVATIVES AND
TITLE OF INVENTION: HEAG PEPTIDES BINDING TO ANNEXINS, THEIR COMPOSITIONS
TITLE OF INVENTION: AND USE
FILE REPREBRUCE: 2551-54
CURRENT APPLICATION NUMBER: US/09/762,270
CURRENT FILING DATE: 2001-02-06
PRIOR APPLICATION NUMBER: PCT/EP99/06231
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
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Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0;
PRIOR APPLICATION NUMBER: PCT/EP99/06231
PRIOR FILING DATE: 1999-08-25
PRIOR APPLICATION NUMBER: EP 98870186.8
PRIOR PLING DATE: 1998-09-01
PRIOR APPLICATION NUMBER: EP 99870062.9
PRIOR FILING DATE: 1999-03-29
NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 22, Application US/09762270; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                               ; ORGANISM: Hepatitis B virus US-09-762-270-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Hepatitis B virus US-09-762-270-22
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US-09-762-270-23
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US-09-762-270-22
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APPLICANT: Samuel E. Hopkins
APPLICANT: Samuel E. Hopkins
APPLICANT: Elizabeth M. Dimassimo
APPLICANT: Bamela P. Rusnak
APPLICANT: Pamela P. Rusnak
TITLE OF INVENTION: Methods and Compositions for Detection of HIV GP41 Peptides and HITLE OF INVENTION: GF41-Like Peptides
TITLE OF INVENTION: GF41-Like Peptides
CURRENT APPLICATION NUMBER: US/09/350,369C
CURRENT APPLICATION NUMBER: 1999-07-09
NUMBER OF SEQ ID NOS: 2041
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Unilever PLC
APPLICANT: Unilever PLC
APPLICANT: Unilever PLC
TITLE OF INVENTION: Peptides Capable of Functioning as Mimotopes for TITLE OF INVENTION: Hormonal Analytes
FILE REFERENCE: Peptide Mimotopes
CURRENT APPLICATION NUMBER: US/09/920,306
CURRENT FILING DATE: 2001-08-02
PRIOR APPLICATION NUMBER: BP03306613.1
PRIOR APPLICATION NUMBER: BP03306613.1
PRIOR PILING DATE: 2000-08-03
NUMBER OF SEQ ID NOS: 80
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic ; OTHER INFORMATION: peptide US-09-920-306-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
65.2%; Score 15; DB 23; Length 5;
Best Local Similarity 100.0%; Pred. No. 4.2e+06;
Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 65.2%; Score 15; DB 23; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                              Query Match 65.2%; Score 15; DB 22; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-350-369C-513
; Sequence 513, Application US/09350369C
; GENERAL INFORMATION:
  ; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-09-841-132-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial Sequence
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US-09-920-306-21
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APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Skeiky, Yasir A.W.
APPLICANT: Probst, Peter
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND
TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION
TITLE OF INVENTION NUMBER: US/09/841,132
CURRENT APPLICATION NUMBER: 2001-04-23
NUMBER OF SEQ ID NOS: 599
SOFTWARE: PSECSEQ for Windows Version 3.0/4.0
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 65.2%; Score 15; DB 22; Length 5; Best Local Similarity 100.0%; Pred. No. 4.2e+06; Matches 2; Conservative 0; Mismatches 0; Indels
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APPLICANT: Guthrie, Kelly
APPLICANT: Guthrie, Kelly
APPLICANT: Merutka, Mohmed
APPLICANT: Anwer, Mohmed
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED
TITLE OF INVENTION: HYBRID POLYPEPTIDES
FILE REFERENCE: 7872-043
CURRENT APPLICATION NUMBER: 09/082,279
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR APPLICATION NUMBER: 09/082,279
PRIOR PILLING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1515
SOFTWARE: PSEUSEQ for Windows Version 3.0
                          PRIOR APPLICATION NUMBER: EP 98870186.8 PRIOR PILING DATE: 1998-09-01 PRIOR APPLICATION NUMBER: EP 99870062.9 PRIOR FILING DATE: 1999-03-29 PRIOR FILING DATE: 1999-03-29 NUMBER OF SEQ ID NOS: 70 SOFTWARE: Patentin Ver. 2.1 SEQ ID NO 23
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; OTHER INFORMATION: Core polypeptide
US-09-834-784-791
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; Sequence 791, Application US/09834784
; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
PRIOR FILING DATE: 1999-08-25
                                                                                                                                                                                                                                                                                     ) ORGANISM: Hepatitis B virus US-09-762-270-23
                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 2; Conserval
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APPLICANT: Samuel E. Hopkins
APPLICANT: Samuel E. Hopkins
APPLICANT: Balizabeth M. Dimassimo
APPLICANT: Toward M. Dimassimo
APPLICANT: Thomas Michael Venetta
APPLICANT: Thomas Michael Venetta
TITLE OF INVENTION: Methods and Compositions for Detection of HIV GP41 Peptides
FILE REFERENCE: 7872-068-999
CURRENT APPLICATION NUMBER: US/09/350,369C
CURRENT FILING DATE: 1999-07-09
NUMBER OF SEQ ID NOS: 2041
SOFTWARE: FastSEQ for Windows Version 4.0
LENGTH: 5
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Sequence 36, Application US/10005305

GENERAL INFORMATION:
APPLICANT: The Government of the United States of America, represented by the APPLICANT: Secretary, Department of Health and Human Services
TITLE OF INVENTION: T20/DP178 AND T21/DP107 ARE ACTIVATORS
TITLE OF INVENTION: OF HUMAN PHAGOCYTE
TITLE OF INVENTION: PRIVAL PEPTIDE RECEPTORS
FILE REFERENCE: NH171.001C1
CURRENT APPLICATION NUMBER: US/10/005,305
CURRENT TILING DATE: 2000-05-05
PRIOR FILING DATE: 2000-06-05
PRIOR FILING DATE: 1999-05-05
NUMBER OF SEQ ID NOS: 203
SEQ ID NOS: 203
LENGTHAE: FASTERE (FWINDOWS Version 4.0)
SEQ ID NO 36
LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT ORGANISM: Human Immunodeficiency Virus-1 (HIV-1)
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                                                                                                                             Sequence 1389, Application US/09350369C GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: PRT
ORGANISM: Artificial Sequence
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1 LW 2
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                                                                           RESULT 74
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PCT-US
217, App
263, App
264, App
94, App
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320, App
34, App
36, App
47, App
47, App
21, App
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Sequence 75, Appl
Sequence 211, App
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(without alignments)
23.274 Million cell updates/sec
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score greater than or equal to the score of the result bein
and is derived by analysis of the total score distribution.
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Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-636-243B-75
US-09-968-74A-211
US-09-968-74A-211
US-09-968-74A-211
US-10-174-717A-94
US-10-174-717A-94
US-10-099-056-2683
US-10-05-33556-34
PCT-US02-33556-34
PCT-US02-33556-34
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PCT-US02-34760-47
PCT-US02-24469A-21
PCT-US02-24469A-21
PCT-US02-24469A-21
PCT-US02-2410-166
PCT-US02-2410-159
PCT-US02-2410-159
PCT-US02-3306-23
PCT-US02-3306-3398
PCT-US02-3410-159
PCT-US02-3410-159
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Maximum Match 100%
Listing first 100 summaries
                                                                                                                            February 12, 2003, 10:51:59
                                                                                         protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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23
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Match Length
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Score

Result No.

61, Appl 26, Appl 14, Appl 114, Appl 115, Appl 1172, Appl 1172, Appl 1172, Appl 1172, Appl 1172, Appl 1173, Appl 1174, Appl 1175, Appl 1175, Appl 1176, Appl 1177, Appl 11

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FILE REPERENCE:

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CURRENT APPLICATION NUMBER: US/09/968,744A

CURRENT FILING DATE:

FRIOR APPLICATION NUMBER: US 9722131.1

PRIOR APPLICATION NUMBER: US 60/065,248

PRIOR PILING DATE: 1997-11-21

PRIOR PILING DATE: 1997-11-21

PRIOR APPLICATION NUMBER: US 60/066,729

PRIOR PILING DATE: 1997-11-21

PRIOR FILING DATE: 1997-11-21

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                                                                                                                                                                                                                 ; Sequence 211, Application US/09968744A; GENERAL INFORMATION: APPLICANT: Tomlinson, Ian M APPLICANT: Winter, Gregory
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 50.05
Matches 1; Conservative
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US-09-968-744A-217
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 4
US-09-968-744A-217
                                                                                                                                                           RESULT 3
US-09-968-744A-211
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APPLICANT: Feng, Weijun
APPLICANT: Feng, Weijun
APPLICANT: Feletterick, Robert
APPLICANT: Rushner, Peter
APPLICANT: West, Brian
APPLICANT: Wammorc, Keith
TITLE OF INVENTION: COACTIVAOR BINDING
FILE REPERENCE: 991-008-999
FILE REPERENCE: 991-008-999
PRIOR PELLING DATE: 1999-03-30
PRIOR APPLICATION NUMBER: US 60/079,956
   Sequence 38, Appl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 82.6%; Score 19; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 5; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
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6 US-10-058-513-38
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US-09-656-243B-75

Sequence 75, Application US/09636243B

GENERAL INFORMATION:

APPLICANT: WANG, Bryan S.

APPLICANT: PADAD, Carl O.

TITLE OF INVENTION: DIMERIZING PEPTIDES

FILE REFERENCE: 8325-1004 / M4-USI

CURRENT FILING DATE: 2000-09-10

PRIOR APPLICATION NUMBER: 05/09/636,243B

CURRENT FILING DATE: 2000-08-10

PRIOR APPLICATION NUMBER: 60/148,422

PRIOR FILING DATE: 1999-08-11

NUMBER OF SEQ ID NOS: 83

SOFTWARE: PATENTIN Ver. 2.0

SEQ ID NO 75
                                                                                              ALIGNMENTS
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; CCATTON: (2)...(3)
: OTHER INPORMATION: Xaa is any amino acid
US-09-281-717A-3
                                                                                                                                                                                                                                                   Sequence 3, Application US/09281717A GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.0
SEQ ID NO 3
LENGTH: 5
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ORGANISM: Artificial Sequence
5
                                                                                                                                                                                                                                                                                                                APPLICANT: Baxter, John
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
47.8
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RESULT 7
US-10-174-717A-94
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                                      SEQ ID NO 2683
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TITLE OF INVENTION: More and more protein-protein interactions in adipocyte cells
FILE REFERENCE: B4838A
CURRENT APPLICATION NUMBER: US/10/099,056
CURRENT FILING DATE: 2002-03-14
PRIOR APPLICATION NUMBER: US 60/275.734
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STREET: 121 South Eighth Street, St. 1600
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 56.5%; Score 13; DB 6; Length 5; Best Local Similarity 50.0%; Pred. No. 3.5e+05; Matches 1; Conservative 1; Mismatches 0; Indels
                                                                                                                                     Sequence 51, Application US/10174717A
APPLICANT: Koide, Shohei
TITLE OF INUNTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: 09/096,749
FILING DATE: June 12, 1998
APPLICATION NUMBER: 60/049,410
FILING DATE: June 12, 1997
ATTORNEY/AGENT INPORMATION:
NAME: Ann S. Viksnins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US4
TELECOMMUNICATION INPORMATION:
TELECHONE: (612) 373-6900
                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: WINDOWS
SOFTWARE: FASTSEQ VECSION 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/174,717A
PRIOR APPLICATION DATA:
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SEQUENCE DESCRIPTION: SEQ ID NO: 51:
US-10-174-717A-51
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GENERAL INFORMATION:
APPLICANT: Hybrigenics
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (612) 339-3
INFORMATION FOR SEQ ID NO: 51:
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                                                                                                                                                                                                                                                                                                                                           ZIP: 55402
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                          USA
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NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, St. 1600
CITY: Minneapolis
                                                                                                                                                                                                                             Score 13; DB 6; Length 5;
Pred. No. 3.5e+05;
1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE: June 12, 1998
APPLICATION NUMBER: 60/049,410
FILING DATE: June 12, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Ann S. Viksnins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US4
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                ; OTHER INFORMATION: Translation of SEQ ID NO:691
US-10-099-056-2683
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: WINDOWS
SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/174,717A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FRAGMENT TYPE: internal ORIGINAL SOURCE: SEQUENCE DESCRIPTION: SEQ ID NO: 94:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: 09/096,749
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 94, Application US/10174717A APPLICANT: Koide, Shohei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE: 18-Jun-2002 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: (612) 373-69
TELEFAX: (612) 339-3061
INFORMATION FOR SEQ ID NO: 94:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                           56.5%;
PRIOR FILING DATE: 2001/03/14
NUMBER OF SEQ ID NOS: 2850
SOFTWARE: PatentIn version 3.1
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MOLECULE TYPE: peptide
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                         Query Match
Best Local Similarity 50.0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
                                                                                                      TYPE: PRT
ORGANISM: Homo Sapiens
FEATURE:
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// Sequence 36, Application PC/TUS0233556
// GENERAL INFORMATION:
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                       Sequence 34, Application PC/TUS0233556 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 100.0
Matches 1; Conservative
                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: PRT
CORGANISM: Homo sapiens
PCT-US02-33556-36
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; ORGANISM: Homo sapiens
PCT-US02-33556-40
PCT-US02-33556-34
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TITLE OF INVENTION: SPECIFIC BINDING AGENTS OF HUMAN ANGIOPOIETIN-2
FILE REFERENCE: A-801B
CURRENT APPLICATION NUMBER: PCT/US02/32657
CURRENT FILING DATE: 2002-10-11
PRIOR APPLICATION NUMBER: US 60/414,155
PRIOR FILING DATE: 2002-99-27
PRIOR PILING DATE: 2001-10-11
NUMBER OF SEQ ID NOS: 359
SOFTWARE: Patentin version 3.1
SEQ ID NO 65
                                                                                                                                                                                                                       APPLICANT: Hoe, Mee H.
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Houghton, Alan
APPLICANT: Ouerfelli, Ouathek
APPLICANT: Moroi, Yoichi
TITLE OF INVENTION: CONJUGATE HEAT SHOCK PROTEIN-BINDING PEPTIDES
FILE REFERENCE: 11746/460018
CURRENT APPLICATION NUMBER: US/10/053,498B
CURRENT FILING DATE: 2002-01-17
PRICA APPLICATION NUMBER: 08/961,707
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 52.2%; Score 12; DB 6; Length 5; Best Local Similarity 40.0%; Pred. No. 3.5e+05; Matches 2; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                         APPLICANT: Sloan-Kettering Institute for Cancer Research APPLICANT: Rothman, James E. APPLICANT: Mayhew, Mark
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Random peptide in coliphage M13
US-10-053-498B-320
                                                                                                              ; Sequence 320, Application US/10053498B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQ ID NOS: 321
SOFTWARE: WordPerfect 8.0 for Windows
SEQ ID NO 320
LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR FILING DATE: 1997-10-31
                                                                                           US-10-053-498B-320
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2 VW 3
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APPLICANT: Amgen, Inc.
TITLE OF INVENTION: Fully Human Antibody Fab Fragments with Human Interferor-Gamma
TITLE OF INVENTION: Neutralizing Activity
FILE REFERENCE: A-799
CURRENT APPLICATION NUMBER: PCT/US02/33556
CURRENT FILING DATE: 2002-10-17
NUMBER OF SEQ ID NOS: 135
SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Amgen, Inc.

TITLE OF INVENTION: Fully Human Antibody Fab Fragments with Human Interferon-Gamma TITLE OF INVENTION: Neutralizing Activity
FILE REFERENCE: A-799
CURRENT APPLICATION NUMBER: PCT/US02/33556
CURRENT APPLICATION NUMBER: PCT/US02/33556
NUMBER OF SEQ ID NOS: 135
SOFTWARE: Patentin version 3.0
SEQ ID NO 36
LENGTH: 5
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TITLE OF INVENTION: Fully Human Antibody Fab Fragments with Human Interferon-Gamma TITLE OF INVENTION: Neutralizing Activity
FILE REPERBNCE: A-799
CURRENT APPLICATION NUMBER: PCT/US02/33556
CURRENT FILING DATE: 2002-10-17
SUPPRENT FILING DATE: 2002-10-17
SOFTWARE: PatentIn version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                        47.8%; Score 11; DB 1; Length 5; 100.0%; Pred. No. 3.5e+05; Live 0; Mismatches 0; Indels
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Best Local Similarity
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PCT-US02-34760-21

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Sequence 47, Application PC/TUS0234760

SEQUENCE 1 INFORMATION:

APPLICANT: Abbott Laboratories

APPLICANT: Bradley, Michael

TITLE OF INVENTION: HAVING ANTIANGIOGENIC ACTIVITY

TITLE OF INVENTION: HAVING ANTIANGIOGENIC ACTIVITY

FILE REPERENCE: 6854, WO. O.

CURRENT APPLICATION NUMBER: PCT/US02/34760

CURRENT PLING DATE: 2002-10-04

PRIOR APPLICATION NUMBER: 10/263,811

PRIOR FILING DATE: 2001-10-14

PRIOR FILING DATE: 2001-10-14

PRIOR PLILING DATE: 2001-10-31

PRIOR APPLICATION NUMBER: 10/000,007

PRIOR FILING DATE: 2001-10-31

NUMBER OF SEQ ID NOS: 61

SEQ ID NO 47
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Bradley, Michael F.
APPLICANT: Bradley, Michael F.
TILLE OF INVENTION: TETRA, PENTA, HEXA- AND HEPTAPEPTIDES
TITLE OF INVENTION: TETRA, PENTA, HEXA- AND HEPTAPEPTIDES
TITLE OF INVENTION: TETRA, PENTANGIOGENIC ACTIVITY
FILE REFERENCE: 6854.WO.01
CURRENT APPLICATION NUMBER: PCT/US02/34760
CURRENT FILING DATE: 2002-10-30
PRIOR APPLICATION NUMBER: 10/23,811
PRIOR PELING DATE: 2001-10-10
PRIOR PELING DATE: 2001-10-31
PRIOR FILING DATE: 2001-10-31
PRIOR FILING DATE: 2001-10-31
PRIOR FILING DATE: 2010-10-31
SPRIOR FILING DATE: 2010-10-31
NUMBER OF SEQ ID NOS: 61
SSOFWWARE: FastSEQ for Windows Version 4.0
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                                                                                     0; Indels
                                          Length 5;
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CTHER INFORMATION: Xaa = proethylamide at position 5
PCT-US02-34760-46
                                        47.8%; Score 11; DB 1; Le 100.0%; Pred. No. 3.5e+05; ative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE: OTHER INFORMATION: Antiangiogenic Peptide
                                                                                                                                                                                                                                                                             PCT-US02-34760-46; Sequence 46, Application PC/TUS0234760; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Abbott Laboratories
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ORGANISM: Artificial Sequence
                                                                                          Conservative
                    Query Match
Best Local Similarity
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; OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids PCT-US02-33985-2
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APPLICANT: Bradley, Michael F.
ITILE OF INVENTION: FEFRA-, PENTA-, HEXA- AND HEPTAPEPTIDES
TITLE OF INVENTION: FORTUNG ANTIANGIOGENIC ACTIVITY
FILE REFERENCE: 6854.WO.01
CURRENT APPLICATION NUMBER: PCT/US02/34760
CURRENT PILING DATE: 2002-10-30
PRIOR FILING DATE: 2002-10-04
PRIOR FILING DATE: 2001-10-31
PRIOR PELICATION NUMBER: 10/000, 540
PRIOR FILING DATE: 2001-10-31
PRIOR PILING DATE: 2001-10-31
PRIOR PILING DATE: 2001-10-31
PRIOR PILING DATE: 2001-10-31
PRIOR PILING DATE: 2001-10-31
NUMBER OF SEQ ID NOS: 61
SOFTWARE: FEASEEG for Windows Version 4.0
SEQ ID NO 21
LENGTH: 5
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OTHER INFORMATION: Xaa = prolylethylamide at position 5
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GENERAL INFORMATION:
APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: 16 Human Secreted Proteins
FILE REFERENCE: PS739PCT
0; Mismatches
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OTHER INFORMATION: Xaa = Nva at position 2
                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: PCT/USO2/33985
CURRENT FILING DATE: 2002-10-24
PRIOR APPLICATION NUMBER: 60/330,629
PRIOR FILING DATE: 2001-10-26
NUMBER OF SEQ ID NOS: 62
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO. 2
LENGTH: 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PCT-US02-34760-21; Sequence 21, Application PC/TUS0234760; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Abbott Laboratories
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Conservative
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ORGANISM: Homo sapiens
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TITLE OF INVENTION: COMPOSITIONS AND METHODS RELATING TO THE DAPTOMYCIN TITLE OF INVENTION: ELOSYNHHETIC GENE CLUSTER FLILE REPERENCE: CUB-12 PCT (IF) CURRENT APPLICATION NUMBER: PCT/US2/24310

CURRENT APPLICATION NUMBER: PCT/US2/24310

CURRENT FILING DATE: 2002-10-25

PRIOR PLICATION NUMBER: PCT/US3354

PRIOR FILING DATE: 2001-10-17

PRIOR FILING DATE: 2001-10-17

PRIOR FILING DATE: 2001-08-06

PRIOR FILING DATE: 2002-05-10

PRIOR FILING DATE: 2002-05-10

PRIOR FILING DATE: 2002-05-10

PRIOR FILING DATE: 2002-05-10

PRIOR FILING DATE: 2002-05-10
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                                                                         47.8%; Score 11; DB 1; Length 5; illarity 100.0%; Pred. No. 3.5e+05; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                           FCT-USO2-24469-21
Sequence 21, Application PC/TUSO224469
GENERAL INFORMATION:
APPLICANT: Merck & Co., Inc.
TITLE OF INVENITON: Biological Assay Detection Method
FILE REFERENCE: PCT 2090'0'
CURRENT APPLICATION NUMBER: PCT/USO2/24469
CURRENT FILING DATE: 2002-08-02
PRIOR FILING DATE: 2001-08-07
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: (1)...(5)
OTHER INFORMATION: Xaa = Any Amino Acid
FEATURE:
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APPLICANT: CUBIST PHARMACEUTICALS, INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: PRT ORGANISM: Artificial Seguence
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SEQ ID NO 166
LENGTH: 5
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Best Local Similarity ?
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NAME/KEY: VARIANT
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       PCT-US02-16525A-26
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LENGTH: 5
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COTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids PCT-US02-35606-2
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47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
TITLE OF INVENTION: Chemokine Beta-1 Fusion Proteins
FILE REPERBNCE: PFSS6PG-1
CURRENT APPLICATION NUMBER: PCT/US02/16525A
CURRENT FILING DATE: 2002-05-24
PRIOR APPLICATION NUMBER: 60/293,212
PRIOR PELLIG DATE: 2001-05-25
NUMBER OF SEQ ID NOS: 137
SOFTWARE: Patentin version 3.1
SEQ ID NO 26
LENGTH: 5
LENGTH: 5
                                                                                                                                                                                                NAME/KEY: VARIANT
LOCATION: (5)...(5)
OTHER INFORMATION: Xaa = proethylamide at position 5
PCT-US02-34760-47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application PC/TUS0235606
| GENERAL INFORMATION:
| APPLICANT: Human Genome Sciences, Inc.
| TITLE OF INVENTION: 41 Human Secreted Proteins
| FILE REFERENCE: PS740FCTTTUS OF UNPER: PCT/US02/35606
| CURRENT FILING DATE: 2002-11-06
| PRIOR APPLICATION NUMBER: 60/331,046
| PRIOR PILING DATE: 2001-11-07
| NUMBER: PETENTION OF PRIOR PLICATION NUMBER: 50/311,046
| WUMBER: PETENTION OF PRIOR PLICATION OF PRIOR PLANCE PLANCE
                                                                                                                                        OTHER INFORMATION: Antiangiogenic Peptide
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                         TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Homo sapiens
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ORGANISM: Homo Sapiens
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Matches 1; Conserva
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LOCATION: (3)
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PCT-US02-35606-2
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LENGTH: 5
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LENGTH: 5
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                                                                                                FEATURE:
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; OTHER INFORMATION: Description of Artificial Sequence:/note; OTHER INFORMATION: Synthetic Construct
PCT-US02-39109-15
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ORGANISM: Murinae gen. sp.
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ORGANISM: Homo sapiens
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PCT-US02-40891-1113
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                                                                                                                                                   Gaps
; OTHER INFORMATION: Description of Artificial Sequence: Illustrative; OTHER INFORMATION: thioesterase motif PCT-US02-24310-166
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                                                                                                 Query Match 47.8%; Score 11; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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1TLLE OF INVENTION: Biological Assay Detection Method
FILE REFERENCE: PCT 20000Y
CURRENT APPLICATION NUMBER: PCT/US02/24469A
CURRENT APPLICATION NUMBER: 60/310,599
PRIOR APPLICATION NUMBER: 60/310,599
PROR PILING DATE: 2001-08-07
NUMBER OF SEQ ID NOS: 27
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 21
LENGTH: 5
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SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 15
LENGTH: 5
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OTHER INFORMATION: Xaa = Any Amino Acid
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COCATION: (1)...(5)
COTHER INFORMATION: Xaa = Any Amino Acid
PCT-US02-24469A-21
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; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
PEATURE:
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Sequence 23, Application PC/TUS0232307
GENERAL INFORMATION:
APPLICANT: IMMUNOMEDICS, INC.
TITLE OF INVENTION: THERAPETIC AGENTS
TITLE OF INVENTION: THERAPETIC AGENTS
TITLE OF INVENTION: THERAPETIC AGENTS
FILE REFERENCE: 018733-1148
CURRENT APPLICATION NUMBER: PCT/US02/32307
CURRENT APPLICATION NUMBER: PCT/US02/32307
CURRENT APPLICATION NUMBER: US 60/416,531
PRIOR FILING DATE: 2002-10-08
NUMBER OF SEQ ID NOS: 27
SOGTWARE: PatentIn version 3.1
SEQ ID NO 23
LENGTH: 5
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; Sequence 5, Application PC/TUS0239643
; GENERAL INFORMATION:
; APPLICANT: Smith, Thomas M
; APPLICANT: Shith, Thomas M
; APPLICANT: Kithey, Terence L
; TITLE OF INVENTION: APPRASE POLYPEPTIDES AND NUCLEIC ACIDS ENCODING SAME
; FILE REFERENCE: 0091830/0502531
; CURRENT APPLICATION NUMBER: PCT/US02/39643
; CURRENT APPLICATION NUMBER: PCT/US02/39643
; NUMBER OF SEQ ID NOS: 11
; SOFTWARE: PatentIn version 3.1
; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 5
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47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
Query Match 47.8%; Score 11; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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FILE REFERENCE: MXI-180PC
CURRENT APPLICATION NUMBER: PCT/US03/00440
CURRENT FILING DATE: 2003-01-07
PRIOR APPLICATION NUMBER: US 60/347649
PRIOR PILING DATE: 2002-01-09
PRIOR APPLICATION NUMBER: US 60/404427
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                      PRIOR FILING DATE: 2002-11-05
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Best Local Similarity
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Matches 1; Conserv
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LENGTH: 5
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OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
APPLICANT: Human Genome Sciences, Inc.
TITLE OF INVENTION: Albumin Fusion Proteins
FILE REPERENCE: PPS.4Pc7
CURRENT APPLICATION NUMBER: PCT/US02/40891
CURRENT FILING DATE: 2002-12-23
FRIOR APPLICATION NUMBER: 60/341,811
FRIOR APPLICATION NUMBER: 60/360,000
FRIOR APPLICATION NUMBER: 60/378,950
FRIOR FILING DATE: 2002-05-10
FRIOR APPLICATION NUMBER: 60/398,008
FRIOR PILING DATE: 2002-05-24
FRIOR APPLICATION NUMBER: 60/411,355
FRIOR PILING DATE: 2002-07-24
FRIOR PILING DATE: 2002-09-18
FRIOR PILING DATE: 2002-10-10
FRIOR PILING DATE: 2002-10-10
FRIOR PILING DATE: 2002-10-11
FRIOR APPLICATION NUMBER: 60/414,984
FRIOR APPLICATION NUMBER: 60/414,611
FRIOR PILING DATE: 2002-10-11
FRIOR PILING DATE: 2002-10-11
FRIOR FILING DATE: 2002-10-13
FRIOR FILING DATE: 2002-10-13
FRIOR FILING DATE: 2002-10-28
FRIOR FILING DATE: 2002-11-05
FRIOR FILING DATE: 2002-01-05
FRIOR FILING DATE: 2002-01-05
FRIOR FILING DATE: 2002-01-05
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47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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FILE REFERENCE: PF5/4PCT
CURRENT APPLICATION NUMBER: PC7/US02/40892
CURRENT FILING DATE: 2002-12-23
PRIOR APPLICATION NUMBER: 60/341,811
PRIOR APPLICATION NUMBER: 60/340,
PRIOR APPLICATION NUMBER: 60/378,950
PRIOR APPLICATION NUMBER: 60/378,950
PRIOR PILING DATE: 2002-05-10
PRIOR PRILING DATE: 2002-05-10
PRIOR PILING DATE: 2002-07-24
PRIOR PILING DATE: 2002-07-24
PRIOR PILING DATE: 2002-07-18
PRIOR PILING DATE: 2002-07-18
PRIOR PILING DATE: 2002-07-18
PRIOR PILING DATE: 2002-07-18
PRIOR PILING DATE: 2002-18
PRIOR PILING DATE: 2002-10-18
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PRIOR APPLICATION NUMBER: 60/420,246
PRIOR ALLING DATE: 2002-10-23
PRIOR APPLICATION NUMBER: 60/423,623
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LOCATION: (3)
OTHER INFORMATION: Xaa equals any of the twenty naturally ocurring L-amino acids
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PRIOR APPLICATION NUMBER: 60/350,358
PRIOR FILING DATE: 2002-01-24
Remaining Prior Application data removed - See File Wrapper or PALM.
WUMBER OF SEQ ID NOS: 858
SOFTWARE: Patentin Ver. 2.0
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APPLICANT: Engelhard, Eric K.
APPLICANT: Engelhard, Eric K.
APPLICANT: Broomers, David W.
TITLE OF INVENTION: NOVEL COMPOSITIONS AND METHODS FOR CANCER FILE REFERENCE: 529452000141
CURRENT APPLICATION NUMBER: PCT/US02/33835
CURRENT FILING DATE: 2002-12-19
PRIOR APPLICATION NUMBER: US 10/004,113
NUMBER OF SEQ 1D NOS: 61
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LOCATION: (3)..(3)
OTHER INFORMATION: "Xaa" at position 3 can be any amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Medarex, Inc. et al. TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES AGAINST CD30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             47.8%; Score 11; DB 1; Length 5; ilarity 100.0%; Pred. No. 3.5e+05; Conservative 0; Mismatches 0; Indels
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Gaps

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RESULT 31
US-08-812-443C-13
US-08-812-443C-13
SEQUENCE 13, Application US/08832443C
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: TSYRLOVA, IRENA
MOLPE, STEPHEN D
TITLE OF INVENTION: INHIBITOR OF STEM CELL PROLIFERATION AND
USES THEREOF
                                                        Query Match

47.8%; Score 11; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
ZIP: 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MS WORD
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/832,443C
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GENERAL INFORMATION:
APPLICANT: GONCZYK, ALFRED
APPLICANT: GONDMAN, SIMON
APPLICANT: DIEFERBACH, BEATE
APPLICANT: FINSINGER, DIRK
APPLICANT: FINSINGER, DIRK
TITLE OF INVENTION: BIOTIN DERIVATIVES
FILE REFERENCE: MERCK-1851
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT FILING DATE: 1997-08-07
PRIOR APPLICATION NUMBER: 08/710,251
PRIOR FILING DATE: 1996-09-13
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APPLICATION NUMBER: US 08/627,173
FILING DATE: 03-APR-1996
INFORMATION FOR SEQ ID NO: 13:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
CITY: ARLINGTON
STATE: VIRGINIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear

MOLECULE TYPE: peptide

SEQUENCE DESCRIPTION: SEQ ID NO: 13:
US-08-832-443C-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 03-Apr-1997
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: <Unknown>
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          PCT-US03-00440-38
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GENERAL INFORMATION:

APPLICANT: Medarex, Inc. et al.

TILLE CANT: Medarex, Inc. et al.

TILLE CANT: MEDALEX, INC. et al.

FILE REPERENCE: MXI-180PC

CURRENT APPLICATION NUMBER: PCT/US03/00440

CURRENT PILING DATE: 2003-01-07

PRIOR APPLICATION NUMBER: US 60/347649

PRIOR FILING DATE: 2002-01-09

PRIOR PILING DATE: 2002-01-09

PRIOR PILING DATE: 2002-08-19

SOFTWARE: PASTECE for Windows Version 4.0

SEQ ID NOS: 53
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JAPPLICANT: MACAIREM, Inc. et al.

TITLE OF INVENTION: HUMAN MONOCLONAL ANTIBODIES AGAINST CD30

FILE REPRENCE: MXI-180PC

CURRENT APPLICATION NUMBER: PCT/US03/00440

CURRENT FILING DATE: 2003-01-07

PRIOR FILING DATE: 2003-01-07

PRIOR FILING DATE: 2002-01-09

PRIOR FILING DATE: 2002-01-09

PRIOR FILING DATE: 2002-01-09

PRIOR FILING DATE: 2002-01-09

PRIOR FILING DATE: 2002-12-06

NUMBER OF SEQ ID NOS: 53

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 38
                                                                                                                                                                                                                                                                                   Query Match 47.8%; Score 11; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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PRIOR FILING DATE: 2002-08-19
PRIOR APPLICATION NUMBER: US 60/431684
PRIOR FILING DATE: 2002-12-06
NUMBER OF SEQ ID NOS: 53
SOFTWARE: PastSEQ for Windows Version 4.0
SEQ ID NO 14
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                                                                                                                                                  LENGTH: 5
TYPE: PRT
ORGANISM: Homo sapiens
PCT-US03-00440-14
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PCT-US03-00440-26
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ORGANISM: Homo sapiens
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TYPE: PRT
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                                                                                                                                                                 OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
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APPLICANT: DIEFENBACH, BEATE
APPLICANT: KESSLER, HORST
APPLICANT: KESSLER, HORST
TITLE OF INVENTION: BIOTIN DERIVATIVES
TITLE REFERENCE: MERCK-1851;
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT FILING DATE: 1997-08-07
FRIOR APPLICATION NUMBER: 08/710,251
PRIOR FILING DATE: 1996-09-13
NUMBER: OF SEQ ID NOS: 175
SEQ ID NO 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: JONCZYK, ALFRED
APPLICANT: GOODMAN, SIMON
APPLICANT: DIFFENDACH, BEATE
APPLICANT: DIFFENDACH, BEATE
APPLICANT: FINSINGER, DIRK
TITLE OF INVENTION: BIOTIN DERIVATIVES
FILE REPERENCE: MERCK-1851
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT FILING DATE: 1997-08-07
PRIOR FILING DATE: 1996-09-13
NUMBER OF SEQ ID NOS: 175
SOFTWARE: PATENTIN VET: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 157, Application US/08908416B
; GENERAL INFORMATION:
; APPLICANT: JONCZYK, ALFRED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 152, Application US/08908416B; GENERAL INFORMATION:
                                                                                             TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
NUMBER OF SEQ ID NOS: 175
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 147
I.FRGTH. 6
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US-08-908-416B-157
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                                                                     ; OTHER INFORMATION: Description of Artificial Sequence: Synthetic; CTHER INFORMATION: peptide
US-08-908-416B-157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
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                                                                                                                                                                                                                  0; Indels
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47.8%; Score 11; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                 Query Match
47.8%; Score 11; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 162, Application US/08908416B
Sequence 162, Application US/08908416B
GENERAL INPORMATION:
APPLICANT: JONCZYK, ALFRED
APPLICANT: GOODMAN, SIMON
APPLICANT: DISENBACH, BEATE
APPLICANT: FINSINGER, HORST
APPLICANT: FINSINGER, HORST
TITLE OF INVENTION: BIOTIN DERIVATIVES
FILE REFREENCE: MERCK-1851
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT APPLICATION NUMBER: US/08/908,416B
NUMBER OF SEQ ID NOS: 175
SOFTWARE: PatentIn Ver: 2.1
SEQ ID NO 162
LENDTH: S
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: GOODMAN, SIMON
APPLICANT: GOODMAN, SIMON
APPLICANT: DIEFENBACH, BEATE
APPLICANT: FINSINGER, HORST
TITLE OF INVENTION: BIOTIN DERIVATIVES
FILE REFERENCE: MERCK-1851
CURRENT APPLICATION NUMBER: US/08/908,416B
CURRENT FILING DATE: 1997-08-07
PRIOR APPLICATION WHERE: 08/710,251
PRIOR APPLICATION NUMBER: 08/710,251
PRIOR APPLICATION NUMBER: 08/710,251
NUMBER OF SEQ ID NOS: 175
SOFTWARE: PATENTIN Ver: 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 167, Application US/08908416B; GENERAL INFORMATION:
APPLICANT: JONCZYK, ALFRED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT ORGANISM: Artificial Sequence
TYPE: PRT
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT ORGANISM: Artificial Sequence
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US-08-908-416B-167
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US-08-908-416B-162
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LENGTH: 5
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APPLICANT: Firestone, Raymond A.
APPLICANT: Freetine, Martin U.
APPLICANT: Lenter, Martin U.
APPLICANT: Lenter, Martin U.
APPLICANT: Lenter, Stefan
APPLICANT: Mack, Juergen
APPLICANT: Mack, Juergen
APPLICANT: Leipert, Dietmar
APPLICANT: Leipert, Dietmar
APPLICANT: Telan, Leila A.
TITLE OF INVENTION: EAP-Activated Anti-Tumour Compounds
FILE REFERENCE: 1/1087 US
CURRENT APPLICATION NUMBER: US/09/533,800
CURRENT APPLICATION NUMBER: US 60/134,136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 47.8%; Score 11; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
     SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
REFERENCE/DOCKET NUMBER: 015389-002610US
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
TELEPAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
                                                                                                                        PRIOR APPLICATION NUMBER: US/08/974,549A
APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19-NOV-1996
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-OCT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,843
                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE: 09-MAY-1997
APPLICATION NUMBER: US 08/911,312
FILING DATE: 14-AUG-1997
                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/721,456
FILING DATE: 22-Nov-2000
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 540:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 16, Application US/09533800
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 540:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR FILING DATE: 1999-05-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JS-09-721-456-540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 39
US-09-533-800-16
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; FEATURE:
; OTHER INFORMATION: Description of Artificial Sequence: Synthetic
; OTHER INFORMATION: peptide
US-08-908-416B-167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: Synthetic OTHER INFORMATION: peptide
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TITLE OF INVENTION: Human Telomerase Catalytic Subunit NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
                                                                                                                             Query Match

47.8%; Score 11; DB 4; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: GOODWAN, SIMON
APPLICANT: GOODWAN, SIMON
APPLICANT: DIFFENBACH, BEATE
APPLICANT: FINSINGER, DIRK
ITILE OF INVENTION: BIOTIN DERIVATIVES
FILE REFERENCE: MERCK-1851;
CURRENT APPLICATION NUMBER: US/08/908,416B;
CURRENT APPLICATION NUMBER: 08/710,251
PRIOR FILING DATE: 1996-09-13
NUMBER OF SEQ ID NOS: 175
SOFTWARE: PATCHIN VET. 2.1
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 38
US-09-721-456-540
; Sequence 540, Application US/09721456
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Morin, Gregg B.
Harley, Calvin B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                           RESULT 37
US-08-908-416B-172
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LENGTH: 5
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Gaps

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GENERAL INFORMATION:

APPLICANT: BOEKE, JEF D.

APPLICANT: BOEKE, JEF D.

APPLICANT: BOEKE, JEF D.

APPLICANT: BOEKE, JEF D.

APPLICANT: BOTHN V.

APPLICANT: BETH, DOMBROSKI A.

TITLE OF INVENTION: COMPOSITIONS AND METHODS OF USE OF MAMMALIAN RETROTRANSPOSONS

FILE REFERENCE: 9596-23U3 (053893-5006-02)

CURRENT APPLICATION NUMBER: US(99/653,8128)

CURRENT FILING DATE: 12000-09-01

PRIOR PILING DATE: 1297-04-28

PRIOR PILING DATE: 1996-11-15

PRIOR FILING DATE: 1996-11-16

NUMBER OF SEQ ID NOS: 152

SOFTWARE: PATENTIN VERSION 3.1

SEQ ID NO 145

LENGTH: S
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APPLICANT: ROZAZIAN. HAIG H.
APPLICANT: BOEKE, JEF D.
APPLICANT: BOEKE, JEF D.
APPLICANT: MCRAN, JOHN V.
APPLICANT: BETH, DOMBROSII A.
TITLE OF INVENTION: COMPOSITIONS AND METHODS OF USE OF MAMMALIAN RETROTRANSPOSONS
TILE REPERBRENCE: 9596-23U3 (055893-5006-02)
CURRENT APPLICATION NUMBER: US/09/653,812B
CURRENT FILING DATE: 1000-09-01
PRIOR PAPLICATION NUMBER: US 08/847,844
PRIOR PAPLICATION NUMBER: US 08/749,805
PRIOR PLING DATE: 1996-11-15
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                                                                                                                                            ; LOCATION: (3)...(3)
; OTHER INFORMATION: Xaa equals any amino acid US-09-726-348B-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 145, Application US/09653812B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 152, Application US/09653812B; GENERAL INFORMATION:
                                                                                          OTHER INFORMATION: WSXWS motif
FEATURE:
NAME/KEY: SITE
                         TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-653-812B-145
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     LENGIH:
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TITLE OF INVENTION: Novel Methods Of Ultrasound Treatment Using Gas Or Gaseous Precur
TITLE OF INVENTION: Filled Compositions
FILE REFERENCE: UNGRIGO
CURRENT APPLICATION WUMBER: US/09/813,484
CURRENT FILING DATE: 2001-03-21
PRIOR APPLICATION NUMBER: 08/929,847
NUMBER OF SEQ ID NOS: 39
SOFTWARE: Patentin version 3.1
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GENERAL INFORMATION:
APPLICANT: Wei, Ying-Fei
TITLE OF INVENTION: Transforming Growth Factor Alpha HIII
TITLE OF INVENTION NUMBER: US/09/726,348B
CURRENT APPLICATION NUMBER: US/09/726,348B
CURRENT APPLICATION NUMBER: 08/778,545
PRIOR APPLICATION NUMBER: 08/778,545
PRIOR PILING DATE: 1997-10-03
PRIOR FILING DATE: 1995-01-03
PRIOR FILING DATE: 1995-12-02
NUMBER: OF SEQ ID NOS: 21
NUMBER: OF SEQ ID NOS: 21
SEQ ID NO 5.
                                                                                                                                                                                                                                    47.8%; Score 11; DB 5; Length 5; ilarity 100.0%; Pred. No. 3.5e+05; Conservative 0; Mismatches 0; Indels
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Query Match

47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: misc_feature
LOCATION: (4)...(4)
COTHER INFORMATION: Xaa is azetidine
US-09-813-484-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-813-484-1; Sequence 1, Application US/09813484; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
NUMBER OF SEQ ID NOS: 28
SOFTWARE: PatentIn version 3.1
                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                OTHER INFORMATION: Precursor
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Best Local Similarity
Matches 1; Conserva
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US-09-726-348B-5
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                                               SEQ ID NO 16
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                                                                                               TYPE: PRT
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APPLICATION NUMBER: US/09/943,120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSES:
STREET: 220 MONIGOMETY Street, Suite 2200
CITY: San Francisco
            .
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                                                                                                                                                                                                                                                      PAPLICANT: Olsen et al.

TITLE OF INVENTION: Human Frezzled-Like Protein
FILLE REFERENCE: PF389C1
CURRENT APPLICATION NUMBER: US/09/610,650B
CURRENT FILING DATE: 2000-06-30
PRIOR PILING DATE: 1998-08-11
PRIOR PILING DATE: 1998-08-11
PRIOR FILING DATE: 1997-08-12
PRIOR FILING DATE: 1998-04-10
PRIOR FILING DATE: 1997-08-12
PRIOR FILING DATE: 1997-08-12
PRIOR FILING DATE: 1998-04-10
NUMBER OF SEQ ID NOS: 27
SOFTWARE: Patentin version 3.1
SQCTWARE: Patentin version 3.1
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47.8%; Score 11; DB 5; Le
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0;
            Mismatches
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APPLICATION NUMBER: US/09/307,579
FILING DATE: 07-May-1999
APPLICATION NUMBER: 08/884,325
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COCATION: (3)...(3)
COTHER INFORMATION: Xaa is any amino acid
US-09-610-650B-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/09943120; GENERAL INFORMATION:
APPLICANT: Rothschild, Kenneth J.
                                                                                                                                                                                                                ; Sequence 18, Application US/09610650B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE: 30-Aug-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
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            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 94104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME/KEY: MISC_FEATURE
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                                                                                                                                                                                               US-09-610-650B-18
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         Matches
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COMPUTER: I BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NATE: US/09/556,972
FILING DATE: 24-Apr-2000
PRIOR APPLICATION NUMBER: US/08/996,338
FILING DATE: 22-DEC-1997
APPLICATION NUMBER: US 74,697/1997
FILING DATE: 12-MRA-1997
APPLICATION NUMBER: UP 215,488/1997
FILING DATE: 28-UU-1997
APPLICATION NUMBER: UP 291,837/1997
FILING DATE: 09-OCT-1997
APPLICATION NUMBER: UP 291,837/1997
FILING DATE: 09-OCT-1997
APPLICATION NUMBER: UP 291,837/1997
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                               Query Match 47.8%; Score 11; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 419 Seventh Street, N.W., Suite 300 CITY: Washington
                                                                                                                                                                                                           ; OTHER INFORMATION: Mutant L1.2mneoI construct US-09-653-812B-152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: BROWDY, ROGER L. REGISTRATION NUMBER: 25,618 REFERENCE/DOCKET NUMBER: TORIGOE=3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULGY: linear
MOLECULG TYPE: peptide
FRAGMENT TYPE: internal fragment
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 44
US-09-556-972-12
| Sequence 12. Application US/09556972
| GENERAL INFORMATION:
| APPLICANT: TORIGOE, Kakuji
| APPLICANT: TORIGOE, Kakuji
| KURIMOTO, Musashi
| TITLE OP INVENTION: POLYPEPTIDES
| UNMBER OF SEQUENCES: 37
| CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: BROWDY AND NEIMARK
PRIOR APPLICATION NUMBER: US 60/006,831
PRIOR FILING DATE: 1995-11-16
NUMBER OF SEQ ID NOS: 152
SOFTWARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: 202-628-5197
                                                                                                                                                               ORGANISM: Artificial Sequence
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SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: D.C.
COUNTRY: USA
ZIP: 20004
                                                                                            SEQ ID NO 152
LENGTH: 5
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ORGANISM: Artificial Sequence
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TITLE OF INVENTION: MICROBIAL BETA-GLUCURONIDASE GENES, GENE PRODUCTS AND TITLE OF INVENTION: USES THEREOF

FILE REFERENCE: 076518-0150

CURRENT APPLICATION NUMBER: 02/09/936,759

CURRENT APPLICATION NUMBER: DO2/100/07107

FRIOR APPLICATION NUMBER: 09/270,957

FRIOR APPLICATION NUMBER: 09/270,957

FRIOR APPLICATION NUMBER: 09/270,957

FRIOR PILING DATE: 1999-03-17

FRIOR FILING DATE: 1999-09-08

FRIOR FILING DATE: 1999-09-08

FRIOR FILING DATE: 1997-09-09

FRIOR FILING DATE: 1997-09-09
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                              NAME: CATTOIL, PETER G.
REGISTRATION NUMBER: 32,837
REGISTRATION NUMBER: 32,837
RELECOMMUNICATION INFORMATION:
TELEPHONE: 415-705-8410
TELEPRAX: 415-397-8338
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                   STRANDEDNESS: not relevant
STRANDEDNESS: not relevant
POPOLOGY: not relevant
MOLECULE TYPE: protein
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-943-120-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: ARIZEKE PHARMACEUTICALS, INC.
APPLICANT: HOUSTON, Lou, L.
APPLICANT: SHERIDAN, Philip, J.
APPLICANT: HAWLEY, Stephen
APPLICANT: GLYNN, Jacqueline, M.
APPLICANT: CHAPIN, Steven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 19, Application US/09969748C GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; Sequence 32, Application US/09936759; GENERAL INFORMATION:
; APPLICANT: JEFFERSON, RICHARD
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
                             ATTORNEY/AGENT INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ) ORGANISM: Escherichia coli
US-09-936-759-32
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RESULT 52
US-09-720-278A-2
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     Query Match
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Sequence 1, Application US/09720278A

GENERAL INFORMATION:

PAPPLICANT: Swart, Pieter Jacob

APPLICANT: Waipers, Maria Elizabeth

APPLICANT: Majjer, Dirk Klaas Fokke

APPLICANT: Hageman, Robert Johan Joseph

APPLICANT: Hageman, Robert Johan Joseph

APPLICANT: Hageman, Robert Johan Joseph

APPLICANT: Wan Den Berg, Jeroen Johannes Maria

TITLE OF INVENTION: Preventing surface infections caused by microorganisms

FILE REFERENCE: 702-00214

CURRENT APPLICATION NUMBER: PS/09/720,278A

CURRENT APPLICATION NUMBER: PS/09/720,278A

FRIOR PRILING DATE: 1999-06-28

PRIOR FILING DATE: 1999-11-06

PRIOR FILING DATE: 1998-11-06

PRIOR FILING DATE: 1998-11-06

PRIOR PRILING DATE: 1998-06-26

PRIOR PRILING DATE: 1998-06-26

NUMBER OF SEQ ID NOS: 28

SOFTWARE: Microsoft Word 97 SR-2

SEQ ID NO 1

LENGTH: 5

LENGTH: DATE: DATE

LENGTH: 5

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                                                                                                                                                                                                 APPLICANT: Rettig, Wolfgang J.
APPLICANT: Lenter, Martin U.
APPLICANT: Lenter, Martin U.
APPLICANT: Beters, Stefan
APPLICANT: Garin-Chea, Pilar
APPLICANT: Leipert, Dietmar
APPLICANT: Leipert, Dietmar
APPLICANT: Leipert, Dietmar
APPLICANT: Telan, Leila A.
TITLE OF INVENTION: FAP-Activated Anti-Tumour Compounds
FILE REFERNCE: 1/1087 US
CURRENT APPLICATION NUMBER: US/09/553,800D
CURRENT FILING DATE: 2000-04-21
PRIOR FILING DATE: 1999-05-14
NUMBER OF SEQ ID NOS: 28
SOOTHARE: PatentIn version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 47.8%; Score 11; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                         Sequence 16, Application US/09553800D GENERAL INFORMATION:
                                                                                                                                                                              APPLICANT: Firestone, Raymond A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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; OTHER INFORMATION: Precursor
US-09-553-800D-16
                                                                                                  US-09-553-800D-16
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LENGTH: 5
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Sequence 2. Application US/09720278A

Sequence 2. Application US/09720278A

GENERAL INFORMATION:

APPLICANT: Swart, Pieter Jacob

APPLICANT: Warth Rujers, Maria Elizabeth

APPLICANT: Majers, Maria Elizabeth

APPLICANT: Hageman, Robert Johan Joseph

APPLICANT: Hageman, Robert Johan Joseph

APPLICANT: Van Den Berg, Jeroen Johannes Maria

TITLE OF INVENTION: Pharmaceutical preparations for use in combatting or

TITLE OF INVENTION: Preventing surface infections caused by microorganisms

FILE REFERENCE: 702-00214

CURRENT FILING DATE: 2001-05-24

PRIOR FILING DATE: 1999-06-28

PRIOR FILING DATE: 1999-11-06

PRIOR PAPLICATION NUMBER: NL 1010284

PRIOR FILING DATE: 1998-11-06

PRIOR FILING DATE: 1998-10-09

SOFTWARE: Microsoft Word 97 SR-2
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APPLICANT: Kuipers, Maria Elizabeth
APPLICANT: Meijer, Dirk Klass Fokke
APPLICANT: Meijer, Dirk Klass Fokke
APPLICANT: Heijer, Dirk Klass Fokke
APPLICANT: Heijer, Dirk Klass Fokke
TITLE OF INVENTION: Preventing surface infections caused by microorganisms
TITLE OF INVENTION: Preventing surface infections caused by microorganisms
FILE REFERENCE: 702-00214
CURRENT APPLICATION NUMBER: US/09/720,278A
CURRENT APPLICATION NUMBER: EP 98203765.7
FRIOR APPLICATION NUMBER: EP 98203765.7
FRIOR APPLICATION NUMBER: EP 98203765.7
FRIOR PILING DATE: 1999-11-06
FRIOR PILING DATE: 1999-11-06
FRIOR PILING DATE: 1999-11-06
FRIOR PILING DATE: 1999-11-06
FRIOR FILING DATE: 1999-11-06
FRIOR FILING DATE: 1999-11-06
FRIOR FILING DATE: 1999-11-06
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Length 5;
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   47.8%; Score 11; DB 5; Le
100.0%; Pred. No. 3.5e+05;
                    100.0%; Pred. nc.
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PRIOR FILING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 28
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                                                                   Conservative
                              Best Local Similarity
Matches 1; Conserv
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PRIOR APPLICATION NUMBER: PCT/EP99/04067
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APPLICANT: Wan Den Berg, Jeroen Johannes Maria
APPLICANT: Van Den Berg, Jeroen Johannes Maria
TITLE OF INVENTION: Pharmaceutical preparations for use in combatting or
TITLE OF INVENTION: Preventing surface infections caused by microorganisms
FILE REFERENCE: 702-002214
CURRENT APPLICATION NUMBER: US/09/720,278A
CURRENT PILING DATE: 1999-06-29
FRIOR FILING DATE: 1999-06-29
FRIOR FILING DATE: 1999-11-06
FRIOR PAPLICATION NUMBER: NL 1010284
FRIOR FILING DATE: 1998-10-09
FRIOR APPLICATION NUMBER: NL 1009505
FRIOR FILING DATE: 1998-10-06
FRIOR FILING DATE: 1998-10-06
FRIOR APPLICATION NUMBER: NL 1009505
FRIOR FILING DATE: 1998-10-06

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APPLICANT: Kuipers, Maria Elizabeth
APPLICANT: Kuipers, Maria Elizabeth
APPLICANT: Mejer, Dirk Klaas Fokke
APPLICANT: Hageman, Robert Johan Joseph
APPLICANT: Van Den Berg, Jercen Johannes Maria
TITLE OF INVENTION: Pherwaccutical preparations for use in combatting or
TITLE OF INVENTION: Preventing surface infections caused by microorganisms
FILE REPREMENCE: 702.002214
CURRENT APPLICATION WORBER: US/09/720,278A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
                                                                                                                                                                                                                                                                                                                                                                                             47.8%; Score 11; DB 5; Length 5; ilarity 100.0%; Pred. No. 3.5e+05; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Swart, Pieter Jacob
APPLICANT: Kuipers, Maria Elizabeth
APPLICANT: Meijer, Dirk Klaas Fokke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 21, Application US/09720278A; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 5, Application US/09720278A GENERAL INFORMATION:
SOFTWARE: Microsoft Word 97 SR-2
SEQ ID NO 3
LENGTH: 5
                                                                                                                                                                  ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity
Matches 1; Conserva
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US-09-720-278A-21
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APPLICANT: Swart, Dieter Jacob
APPLICANT: Wart, Dieter Jacob
APPLICANT: Wanter, Maria Elizabeth
APPLICANT: Wipers, Maria Elizabeth
APPLICANT: Wipers, Maria Elizabeth
APPLICANT: Webert Johan Joseph
APPLICANT: Hageman, Robert Johan Joseph
APPLICANT: Van Den Berg, Jeroen Johannes Maria
APPLICANT: Van Den Berg, Jeroen Johannes Maria
TITLE OF INVENTION: Preventing surface infections caused by microorganisms
FILE REFERENCE: 702-002214
CURRENT PELLING DATE: 2001-05-24
PRIOR APPLICATION NUMBER: BP 98203765.7
PRIOR APPLICATION NUMBER: BP 98203765.7
PRIOR PELLING DATE: 1998-11-06
PRIOR PLING DATE: 1998-11-06
PRIOR PLING DATE: 1998-10-09
PRIOR PLING DATE: 1998-10-09
PRIOR PLING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 28
SOFTWARRE: Microsoft Word 97 SR-2
SECTION 23
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PRIOR FILING DATE: 1999-06-28
PRIOR APPLICATION NUMBER: EP 98203765.7
PRIOR APPLICATION NUMBER: BP 98203765.7
PRIOR APPLICATION NUMBER: NL 1010284
PRIOR FILING DATE: 1998-10-09
PRIOR FILING DATE: 1998-06-26
NUMBER OF SEQ ID NOS: 28
SOSTWARE: Microsoft Word 97 SR-2
SEQ ID NO 21
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Matches 1; Conservative
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US-09-636-243B-44
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## APPLICANT: Kuipers, Maria Elizabeth
## APPLICANT: Meijer, Dirk Klaas Fokke
## APPLICANT: Wan Den Berg, Jercen Johannes Maria
## TITLE OF INVENTION: Preventing Surface infections caused by microorganisms
## TITLE OF INVENTION: Preventing surface infections caused by microorganisms
## TITLE OF INVENTION: Preventing Surface infections caused by microorganisms
## TITLE OF INVENTION: Pharmaceutical preparations caused by microorganisms
## TITLE OF INVENTION: 2001-05-24
## PRIOR PILING DATE: 1999-06-28
## PRIOR APPLICATION NUMBER: EP 98203765.7
## PRIOR APPLICATION NUMBER: BP 98203765.7
## PRIOR APPLICATION NUMBER: M. 1010284
## PRIOR PILING DATE: 1998-10-09
## PRIOR FILING DATE: 1998-06-26
## NUMBER OF SEQ ID NGS: 28
## SOFTWARE: Microsoft Word 97 SR-2
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; OTHER INFORMATION: Description of Artificial Sequence: sequential
; OTHER INFORMATION: block reoptimization sequence
US-09-636-243B-30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 30, Application US/09636243B
GENERAL INFORMATION:
APPLICANT: WANG, BTYS
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-USI
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT APPLICATION NUMBER: 05/04/84
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
PRIOR FILING DATE: 1999-08-11
SOFTWARE: Patentin Ver. 2.0
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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US-09-636-243B-30
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OTHER INFORMATION: Description of Artificial Sequence: sequential
COTHER INFORMATION: block reoptimization sequence
US-09-636-243B-35
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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GENERAL INFORMATION:
APPLICANT: WANG, Bryan S.;
APPLICANT: WANG, Bryan S.;
TITLE OF INVENTION: DIMERIZING PREPTIDES
FILE REFERENCE: 8325-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B;
CURRENT FILING DATE: 2000-08-10
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 44
          TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN VEY: 2.0
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US-09-636-243B-56
; Sequence 56, Application US/09636243B
; GENERAL INFORMATION:
; APPLICANT: WANG, Bryan S.
; APPLICANT: PABO, Carl O.
; TITLE OF INVENTION: DIMERIZING PEPTIDES
; FILE REFERENCE: 8325-1004 / M4-US.
; CURRENT APPLICATION NUMBER: US/09/636,243B
; CURRENT FILING DATE: 2000-08-10
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PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN VEY: 2.0
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ORGANISM: Artificial Sequence
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PABO, Carl O.
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; OTHER INFORMATION: Description of Artificial Sequence: sequential
; OTHER INFORMATION: block reoptimization sequence
US-09-636-243B-58
                                                                                                       OTHER INFORMATION: Description of Artificial Sequence: sequential OTHER INFORMATION: block reoptimization sequence
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47.8%; Score 11; DB 5; Le
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 57, Application US/09636243B
GENERAL INFORMATION:
APPLICANT: WANG, Bryan S.;
APPLICANT: WANG, Bryan S.;
APPLICANT: PABO, Carl O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REPERENCE: 8325-1004 / M4-USI
CURRENT PELING DATE: 2000-08-10
PRIOR PEPLICATION NUMBER: US/09/636,243B
CURRENT PILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATCENTIN VOICE: 2.0
SEQ ID NO 57
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APPLICANT: WANG, Bryan S.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8225-1004 / M4-USI
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN Ver. 2.0
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                                        TYPE: PRT ORGANISM: Artificial Sequence
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SEQ ID NO 56
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Query Match 47.8%; Score 11; DB 5; Length 5; Best Local Similarity 100.0%; Pred. No. 3.5e+05; Matches 1; Conservative 0; Mismatches 0; Indels
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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GENERAL INFORMATION:
APPLICANT: WANG, BYAN S.
ATTLE OF INVENTION: DIMERIZING PEPTIDES
FILE REPERRENCE: 8125-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTING VET. 2.0
SEQ ID NO 60
SEQ ID NO 60
                                                                                                                                                                                                                                                                                                                        APPLICANT: WANG, BEYAN S.
APPLICANT: PABO, Carl O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 59
LENGTH: 5
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US-09-636-243B-60
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PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: sequential
OTHER INFORMATION: block reoptimization sequence
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
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                                                                   APPLICANT: WANG, Eryan S.
APPLICANT: WANG, Eryan S.
APPLICANT: PABO, CALI O.
TILE REFERENCE: 8325-1004 / M4-USI
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN Ver. 2.0
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APPLICANT: PABO, Carl O.
TITLE OF INVENTION: DIMERIZING PEPTIDES.
TITLE REFERENCE: 8325-1004 / M4-US.
CURRENT APPLICATION NUMBER: US/09/636,243B; CURRENT FILING DATE: 2000-08-10
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN Ver. 2.0
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APPLICANT: WANG, Carl O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REPERENCE: 8125-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
US-09-636-243B-61; Sequence 61, Application US/09636243B; GENERAL INFORMATION:
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LENGTH: 5
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OTHER INFORMATION: Description of Artificial Sequence: sequential

OTHER INFORMATION: block reoptimization sequence
US-09-636-243B-64
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US-09-636-243B-64
; Sequence 64, Application US/09636243B
; GENERAL INFORMATION:
; APPLICANT: WANG, Bryan S.
; TITLE OF INVENTION: DIMERIZING PEPTIDES
; TITLE OF INVENTION: DIMERIZING PEPTIDES
; CURRENT APPLICATION NUMBER: US/09/636,243B
; CURRENT FILING DATE: 2000-08-10
; PRIOR FILING DATE: 1999-08-11
; NUMBER OF SEQ ID NOS: 83
; SOFTWARE: PATENTIN UNCES: 2000-08-10
; NUMBER OF SEQ ID NOS: 83
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GENERAL INFORMATION:
APPLICANT: WANG, Bryan S.
APPLICANT: WANG, Bryan S.
APPLICANT: WANG, Bryan S.
FILE PERERENCE: 8125-1004 / M4-US1
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR APPLICATION WUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN Ver. 2.0
                      PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 63
LENGTH: 5
CURRENT FILING DATE: 2000-08-10
                                                                                                                                                                                TYPE: PRT
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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US-09-636-243B-76
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APPLICANT: MAEDA, Hiroaki,
APPLICANT: WAEDA, Hiroaki,
APPLICANT: WSHIO, Yoshitaka
APPLICANT: USHIO, Yoshitaka
APPLICANT: USHIO, Yoshitaka
APPLICANT: HIGUCHI, Hirotumi
APPLICANT: HIGUCHI, Hirotumi
APPLICANT: HIGUCHI, Hirotumi
APPLICANT: HIGUCHI, Himminized Immunoglobulins Specifically Reactive to Fas Ligand or
TITLE OF INVENTION: Fragments Thereof, and Apoptosis-Induced Site From Fas Ligand
FILE REFERENCE: 050006-0055
CURRENT APPLICATION NUMBER: US/09/254,180C
CURRENT APPLICATION NUMBER: CT/0P97/02983
FRIOR APPLICATION NUMBER: 271546/1996
FRIOR FILING DATE: 1996-09-20
FRIOR APPLICATION NUMBER: 231472/1996
FRIOR FILING DATE: 1996-09-02
FRIOR APPLICATION NUMBER: 231472/1996
FRIOR FILING DATE: 1996-09-02
NUMBER OF SEQ ID NOS: 183
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APPLICANT: Resqualini, Renata
APPLICANT: Resqualini, Renata
APPLICANT: Rajotte, Daniel
TITLE OF INVENTION: Methods of Identifying Lung Homing Molecules Using
TITLE OF INVENTION: Membrane Dipeptidase
FILE REFERENCE: P-LA 4377
CURRENT PELICATION NUMBER: US/09/676,475A
CURRENT FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 452
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 278
LENGTH: 5
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ORGANISM: Artificial Sequence
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APPLICANT: EDA, Yasuvu
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Best Local Similarity
Matches 1; Conserva
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ORGANISM: Mouse
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                                                                                                                           RESULT 73
US-09-254-180C-1
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                     OTHER INFORMATION: Description of Artificial Sequence: sequential; OTHER INFORMATION: block reoptimization sequence
US-09-636-243B-76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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47.8%; Score 11; DB 5; Length 5;
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                              0; Indels
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                                                                                                                         Query Match
47.8%; Score 11; DB 5; Le
Best Local Similarity 100.0%; Pred. No. 3.5e+05;
Matches 1; Conservative 0; Mismatches 0;
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GENERAL INPORMATION:
APPLICANT: WANG, Bryan S.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERRICE: 8325-1004 / M4-US1
CURRENT PAPLICANTE: 2000-08-10
PRIOR APPLICATION NUMBER: 06/148,422
PRIOR PLING DATE: 1999-08-11
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR FILING DATE: 1999-08-11
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PATENTIN VET. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: WANG, Bryan S.
APPLICANT: PABO, Carl O.
TITLE OF INVENTION: DIMERIZING PEPTIDES
FILE REFERENCE: 8325-1004 / M4-USI
CURRENT APPLICATION NUMBER: US/09/636,243B
CURRENT FILING DATE: 2000-08-10
PRIOR APPLICATION NUMBER: 60/148,422
PRIOR APPLICATION NUMBER: 60/148,422
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 77, Application US/09636243B; GENERAL INFORMATION:
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US-09-636-243B-77
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FEATURE:
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RESULT 75
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Sequence 1, Application US/09920262A
; GENERAL INFORMATION:
    APPLICANT: Seallon, David
    APPLICANT: Might, David
    APPLICANT: Might, David
    APPLICANT: Might, David
    APPLICANT: Might, David
    APPLICANT: Giles Komar, Jull
    APPLICANT: Peritt, David
    APPLICANT: Peritt, David
    APPLICANT: Peritt, David
    APPLICANT APPLICATION NUMBER: US/09/20,262A
    CURRENT FILING DATE: 2002-06-06
    PRIOR APPLICATION NUMBER: 60/223,386
    PRIOR APPLICATION NUMBER: 60/236,827
    PRIOR APP
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103, App 5171838

Sequence 103,

Patent No.

Sequence Sequence

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Sequence 228, App Sequence 148, App Sequence 47, Appl

Sequence

Title: Perfect score:

Sequence:

protein

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Run on:

Scoring table:

Searched:

Database

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US-08-477-362-83
US-08-477-462-83
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US-08-1313-02-4
US-08-1313-02-4
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Sequence 5, Appli
Sequence 5, Appli
Sequence 3, Appli
Sequence 37, Appli
Sequence 44, Appl
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Sequence 11
Sequence 10
Sequence 44
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/cgn2_6/ptodata/1/iaa/RecomB.pep:*
/cgn2_6/ptodata/1/iaa/PcTUS_COMB.pep:*
             GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-424-957-5
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US-09-035-686-4
US-09-035-686-4
US-08-073-028-44
US-08-073-028-44
US-08-133-011-148
US-08-133-011-128-13
US-08-131-122-15
US-08-131-122-15
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US-08-131-132-148
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Maximum Match 100%
Listing first 100 summaries
                                              - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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Match Length
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Sequence 19, Appl Sequence 37, Appl Sequence 51, Appl Sequence 72, Appl Sequence 26, Appl Sequence 11, Appl Sequence 11, Appl Sequence 12, Appl Sequence 9, Appl Patent No. 5217869 Patent No. 5217869 Patent No. 5217869

Sequence Sequence Sequence

Sequence 4, Appli Sequence 3, Appli

Sequence 83, Appl Sequence 83, Appl Sequence 83, Appl Sequence 9, Appli Sequence 1, Appli Sequence 19, Appl Sequence 10, Appl Sequence 10, Appl Sequence 11, Appl Sequence 12, Appl Sequence 5, Appli Sequence 5, Appli Sequence 5, Appli Sequence 2, Appli Sequence 31, Appli Sequence 37, Appli Sequence

Sequence Seq

Sequence 39, Appl Sequence 19, Appl Sequence 21, Appl Sequence 21, Appl Sequence 39, Appl Sequence 31, Appli Sequence 31, Appli

Sequence 15, Sequence 16, Sequence 17,

Sequence

Score

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                                                                                                                          Sequence 4, Application US/08424957
Patent No. 5770377
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSER: Flahr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:

GENERAL INFORMATION:

APPLICANT: Picksley, Steven M.

APPLICANT: Picksley, Steven M.

APPLICANT: Lane, David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:

ADDRESSER: Plehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400

CITY: San Prancisco
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82.6%; Score 19; DB 1; Length 5;
Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 5; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: California
COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IB PC compatible
COMPUTER: IB PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-UL-1994
ATTORNEY/AGENT INFORMATION:
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Patent No. 5770377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELERA: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 4:
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LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: unknown
US-08-424-957-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 FXXLW 5
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Sequence 4, Application US/09035686
Patent No. 6153391
GENERAL INFORMATION:
APPLICANT: Dicksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          82.6%; Score 19; DB 1; Length 5; 80.0%; Pred. No. 2e+05; tive 0; Mismatches 1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400 CITY: San Francisco STATE: California COUNTRY: United States ZIP: 94111-4187
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                 A-61228/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            19-APR-1995
TUMBER: US 08/277,660
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                          NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-611
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 398-3249
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/4
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/2
FILING DATE: 20-UUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
                                                                                                                                                                                                                                                                                                                                                                                             TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 80.v.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: unknown US-08-424-957-5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 FXXLW 5
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US-09-053-3
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Sequence 5, Application US/09035686

Fatent No. 615391

GENERAL INFORMATION:

APPLICANT: Lane, David P.

ITILE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof

CORRESPONDENCES: 50

CORRESPONDENCE ADDRESS:

ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert

STREET: Four Embarcadero Center, Suite 3400

CITY: San Francisco

STATE: California

COUNTRY: United States

ZIP: 9411-4187

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 82.6%; Score 19; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 5; Conservative 0; Mismatches 0; Indels
| REGISTRATION NUMBER: 24,190 | REFERENCE/DOCKET NUMBER: A-61228/WHD | TELECOMMUNICATION INFORMATION: | TELEPAX: (415) 398-3249 | TELEPAX: (415) 398-3249 | INFORMATION FOR SEQ ID NO: 4: | IN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFENCE/DOCKET NUMBER: A-61228/WHD
FELECOMMUNICATION INFORMATION:
TELEPRAN: (415) 781-1989
TELEPRA: (415) 398-3249
TELEPRA: 910 277299
INFORMATION FOR SEQ ID NO: 5:
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APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-UUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 80.0
Matches 4; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                            ; STRANDEDNESS:
; TOPOLOGY: unknown
US-09-035-686-4
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US-09-035-686-5
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CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 FXXLW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTHER INFORMATION: "Xaa" in position 3 stands for OTHER INFORMATION: either Glu or Asp. US-09-099-053-3
Sequence 3, Application US/09099053
Patent No. 6388063
GENERAL INFORMATION:
APPLICANT: Greg Plowman
APPLICANT: Susan Onrust
APPLICANT: Bara Courtneidge
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF
TITLE OF INVENTION: DIAGNOSIS AND TREATMENT OF
TITLE OF INVENTION: SAD RELATED DISORDERS
NUMBER OF SEQUENCES: 28
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 69.6%; Score 16; DB 4; Best Local Similarity 60.0%; Pred. No. 2e+05; Matches 3; Conservative 1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPOTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FEASEEG for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/099,053
FILING DATE: HETEWITh
CLASSIFICATION DATA:
RRICK APPLICATION DATA:
APPLICATION NUMBER: 60/049,914
FILING DATE: June 18, 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                           ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 35" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 235/121
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 6
US-07-893-930-37
is Sequence 37, Application US/07893930
is Patent No. 5317014
is GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Warburg, Richard J. REGISTRATION NUMBER: 32,327
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5 amino acids
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                                                                                                                                                                                                                                                                                        STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: STRANDEDNESS: BILL
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0; Gaps
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           Kucharczyk, Nathalie
Morris, Angela D.
Paladino, Joseph
Bonnet, Joseph
Thurieau, Christophe
WENTION: NEW PEPTIDES AND PSEUDO-
WENTION: COMPOUNDS DERIVED FROM TACHYKININ
                                                                                                                                                                                    NUMBER OF SEQUENCES: 46
CORRESPONDENCE ADDRESS:
ADDRESSEE: Gordon W. Hueschen
STREET: 715 The "H" Bldg., 310 East Michigan
                                                                                                                                                                                                                                                                                                                                                                COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: #1.25
SOFTWARE: #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/893,930
FILING DATE: 19920603
CLASSIFICATION ATA:
APPLICATION NUMBER: FR 91.06721
FILING DATE: 19920603
CLASSIFICATION DATA:
APPLICATION NUMBER: FR 91.06721
FILING DATE: 19920603
FILING APPLICATION NUMBER: FR 91.06721
FILING DATE: 16.157
REFERENCE/DOCKET NUMBER: 16,157
TELECOMMUNICATION INFORMATION:
TELEFRANCE CLIG-382-0030
TELEFRANCE CLIG-382-2030
TELEFRANCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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OTHER INFORMATION: /note= "Benzhydryl
OTHER INFORMATION: carbonyl"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: /note= "NH2"
US-07-893-930-37
Fauchere, Jean-Luc
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                                                                           APPLICANT: BODDET, J. APPLICANT: Thurieau, TITLE OF INVENTION: TITLE OF INVENTION: TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: AMINO ACID
TOPOLOGY: linear
MOLECULE TYPE: mRNA
                                                                                                                                                                                                                                                                                                  Kalamazoo
                                                                                                                                                                                                                                                                                                                                                USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 7
US-08-073-028-44
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APPLICANT:
                  APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                             COUNTRY:
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                                                             APPLICANT
                                                                                                                                                                                                                                                                                                                      STATE:
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; Sequence 44, Application US/08073028
; Patent Mo. 5464933
; GENERAL INFORMATION:

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0; Gaps
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| Patent No. 5658727|
| Patent No. 5658727|
| Patent No. 5658727|
| APPLICANT: Rang, Angray
| APPLICANT: Barbas, Carlos
| APPLICANT: Lerner, Richard A. |
| TITLE OF INVENTION: HETERODIMERIC RECEPTOR LIBRARIES USING |
| TITLE OF INVENTION: HETERODIMERIC RECEPTOR LIBRARIES USING |
| TITLE OF INVENTION: HETERODIMERIC RECEPTOR LIBRARIES USING |
| TITLE OF INVENTION: HETERODIMERIC RECEPTOR LIBRARIES USING |
| TITLE OF INVENTION: HETERODIMERIC RECEPTOR LIBRARIES USING |
| TITLE OF INVENTION: HETERODIMERIC OF ADDRESSEE: The Scripps Research Institute, Office of ADDRESSEE: Patent Counsel; ADDRESSEE: Patent Counsel; STREET: 10666 No. 5658727th Torrey Pines Road, TPC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 65.2%; Score 15; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                       TITLE OF INVENTION: SYNTHETIC PEPTIDE INHIBITORS OF HIV TITLE OF INVENTION: TRANSMISSION NUMBER OF SEQUENCES: 74
CORRESPONDENCE ADDRESS: ADDRESSE: Pennie & Edmonds STREET: 1155 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER TOOS AND TOO TO THE TOO THE TO
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/133,011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
APPLICANT: Bolognesi, Dani P. APPLICANT: Matthews, Thomas J. APPLICANT: Wild, Carl T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          5 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 8
US-08-133-011-148
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SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                           LENGTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide;
HYPOTHETICAL: YES
US-08-584-579-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              , MOLECULE TYPE: peptide US-08-704-170-103
       PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
GY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-704-170-103
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; Patent No. 5674979
; GENERAL INFORMATION: APPLICANT: Schramm, Wolfgang
APPLICANT: Schramm, Hans J.
; TITLE OF INVENTION: Agent For Inhibiting Symmetrical
TITLE OF INVENTION: Proteins, In Particular Enzymes
NUMBER OF SEQUENCES: 20
; CORRESPONDENCE ADDRESS:
ADDRESSEE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          65.2%; Score 15; DB 1; Length 5; 100.0%; Pred. No. 2e+05; tive 0; Mismatches 0; Indels
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ZIP: 20005

MEDIUM TYPE: Floppy disk
COMPUTER: EM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,579
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/332,447
FILING DATE: 31-0CT-199
APPLICATION NUMBER: US 08/112,215
FILING DATE: 26-AUG-1993
APPLICATION NUMBER: US 07/976,003
FILING DATE: 13-NOV-1992
APPLICATION NUMBER: US 07/585,141
FILING DATE: 07-DEC-1990
APPLICATION NUMBER: WO PCT/EP90/00219
FILING DATE: 09-FEB-1990
         PRIOR APPLICATION 1435
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: US 07/683,602
FILING DATE: 10-APR-1991
PRIOR APPLICATION NUMBER: US 07/826,623
FILING DATE: 27-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US 92/03091
FILING DATE: 10-APR-1992
ATTORNEY AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: SCRF 238.2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 619-554-6312
INFORMATION FOR SEQ ID NO: 148:
SEQUENCE CHARACTERISTICS:
08-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: protein FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity
Matches 2; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
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FILMS DATE: WINDERS 139 94 040.2

FILMS DATE: WINDERS 12997

ATTORNY CREAT INCOMPTION: WINDERS 12997

**RESTREMENDERS 12997

**RESTREMEND
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APPLICANT: HILGER, CHRISTOPH-STEPH
APPLICANT: DINKELBORG, LUDGER
APPLICANT: KRAMP, WOLGANG
APPLICANT: SCHIER, HANS-WARTIN
ITITLE OF INVENTION: CHELATORS OF TYPE XNISIOI FOR RADIOACTIVE ISOTOPES, 7H
NUMBER OF SEQUENCES: 15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               65.2%; Score 15; DB 2; Length 5; 100.0%; Pred. No. 2e+05; tive 0; Mismatches 0; Indels
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Patent No. 5562642
GENERAL INFORMATION:
APPLICANT: TSUGITA, Akira
TITLE OF INVENTION: Method for Sequencing of
TITLE OF INVENTION: Protein or Peptide
NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Adams & Wilks
STREET: 50 Broadway, 31st Floor
CINTER NOW YOUR.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REGISTRATION NUMBER: 25,329
REPERENCE/DOCKET VUMBER: WBLT 2006 (WO) PA
TELECOMMUNICATION:
TELEPHONE: 415-362-3800
                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: FLIESLER, DUBB, MEYER & LOVEJOY STREET: Four Embarcadero Center, Suite 400 CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Pred. wc.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              UMBER: P 43 10 999.3
31-MAR-1993
                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM: DOS
SOFTWARE: FASTSEQ VERSION 1.5
SOFTWARE: APPLICATION DATA:
APPLICATION NUMBER: US/08/530,326
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      14:
                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible OPERATING SYSTEM: DOS SOFTWARE: FastSEQ Version
                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/D
FILING DATE: 29-MAR-1994
APPLICATION NUMBER: P 43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 31-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Dubb, Hubert E
REGISTRATION UMBER: 25,3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: 415-362-380
TELEFAX: 415-362-2928
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MEDIUM TYPE: Diskett
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Best Local Similarity
Matches 2; Conservat
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    GENERAL INFORMATION:
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US-08-854-187A-2
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COUNTRY:
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                                                                                                                                                                                                                                              COUNTRY:
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TITLE OF INVENTION: NON-CROSSLINKED PROTEIN PARTICLES FOR
TITLE OF INVENTION: THERAPEUTIC AND DIAGNOSTIC USE
NUMBER OF SEQUENCES: 184
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 2; Length 5; larity 100.0%; Pred. No. 2e+05; Conservative 0; Mismatches 0; Indels
Query Match 65.2%; Score 15; DB 1; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Felease #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/747,137 FILING DATE: 12-NOV-1996 CLASSIFICATION: 424 PRIOR APPLICATION DATA: PAPLICATION NUMBER: US/08/212,546 FILING DATE: 14 PAR-1994 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     016197-000840US
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FILING DATE: 01-JUN-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/959,560
FILING DATE: 13-OCT-1992
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US 07/641,720
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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US-08-530-336-14
; Sequence 14, Application US/08530326
; Patent No. 5961954
                                                                                                                                                                                                                        Sequence 44, Application US/08747137
Patent No. 5945033
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Apple, Randolph T.
REGISTRATION UNDHER: 36,429
REFERENCE, DOCKET UNDHER: 01615
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-576-0200
INFORMATION FOR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US 0
FILING DATE: 15-JAN-1991
ATTORNEY/AGENT INFORMATION:
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Matches 2, Conserva
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INFORMATION FOR SEQ ID NO:
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US-08-862-625A-3
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TITLE OF INVENTION: MONOSPECIFIC ANTIBODY REACTIVE WITH
TITLE OF INVENTION: MATRIX METALLOPROTEINASE CLEAVAGE PRODUCTS OF FIBRIN(OGEN)
NUMBER OF SEQUENCES:
ADDRESSEE: Hoffmann & Baron, LLP
STREET: 350 Jericho Turnpike
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 65.2%; Score 15; DB 2; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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MEDIUM TYPE: Diskette, 3.50 inch, 1.44 Mb storage COMPUTER: IBM compatible
OPERATING SYSTEM: MS-DOS
                 COMPUTER READBLE FORM:

MEDIUM TYPE: Disketre, 3.50 inch,
MEDIUM TYPE: 1.44 Mb storage
COMPUTER: NEC PowerMate VP75
CORDUTER: Wordperfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/854,187A
FILING DATE: 09-MAY-1997
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: JP 08-115087
FILING DATE: 09-MAY-1996
APPLICATION NUMBER: JP 08-154580
FILING DATE: 19-4-JW1-1996
APPLICATION NUMBER: JP 08-154580
FILING DATE: 14-JW1-1996
APPLICATION NUMBER: JS,386
REFERENCE/DOCKET NUMBER: S5,386
REFERENCE/DOCKET NUMBER: S5,386
REFERENCE/DOCKET NUMBER: S5,386
REFERENCE/LOCKET NUMBER: S5,386
REFE
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CURRENT APPLICATION DATA:
APPLICATION NDRAR:
TLING DATE: 7/25/97
CLASSIFICATION: 435
ATTORNEY AGANT INFORMATION:
NAME: 0'Dea, Sean W.
REGISTRATION NUMBER: 37690
REGISTRATION NUMBER: 454-16
TELECHONE: (516) 822-3550
TELEFAX: (516) 822-3550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/08900895
Patent No. 6043087
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: not applicable
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: linear; MOLECULE TYPE: peptide US-08-854-187A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Jericho
STATE: New York
COUNTRY: USA
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PROGRAMIN SEGULATION:

TYPE: Annua scaled state of the segulation scale of the
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OTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION NUMBER: US/08/981,1228
CURRENT FILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: UP 7-176904
PRIOR FILING DATE: 1995-06-11
PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR FILING DATE: 1996-06-21
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; Sequence 51, Application US/08981122B
; Patent No. 6127339
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ORGANISM: Artificial Sequence
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NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILE REFERENCE:
          ; OTHER INFORM
US-08-981-122-38
                                                                                                                                                                                                                                                                                                                  US-08-981-122-43
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SEQ ID NO 51
LENGTH: 5
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LENGTH: 5
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OTHER INFORMATION: Sequence of a peptide synthesized in Examples 11 and 12 from L-fd
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Patent No. 6127339
OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Patent No. 6127339
OTHER INFORMATION: F-moc amino acids by solid phase method using a multipeptide
OTHER INFORMATION: synthesizing system (RaMPS)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FEATURE:
OTHER INFORMATION: Sequence of a peptide synthesized in Example 2 from L-form
                                                   0;
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                                                                                                                                                                                                                                                                                                                                    APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Azitomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/08/981,122B
CURRENT PILING DATE: 1997-12-18
PRIOR FILING DATE: 1997-62-1
PRIOR FILING DATE: 1996-06-21
PRIOR FILING DATE: 1996-06-21
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APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Gaps
                                                   0; Gaps
Query Match 65.2%; Score 15; DB 3; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 2; Conservative 0; Mismatches 0; Indels
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CURRENT FILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: US/08/981,122B
PRIOR PILING DATE: 1995-06-21
PRIOR APPLICATION NUMBER: PT/JP96/01734
PRIOR PILING DATE: 1996-06-21
PRIOR PILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: PATENTIN VET: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 38, Application US/08981122B Patent No. 6127339
                                                                                                                                                                                                                                                               ; Sequence 15, Application US/08981122B ; Patent No. 6127339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
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SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY: AMIDATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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LENGTH: 5
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NAME/KEY: AMIDATION
LOCATION: 5
COCATION: 5
OTHER INFORMATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for Patent No. 6127339
OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid OTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
                                                                           0
                                                                                                                                                                                                                                                                                                                                                                            Sequence 43, Application US/08981122B
Patent No. 6127339
GENERAL INFORMATION:
APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Aricomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Hatenaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Gaps
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                                                                    0; Indels
Query Match 65.2%; Score 15; DB 3; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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NAME: Mortimer, John S.
REGISTRATION NUMBER: 30,407
REFERENCE/DOCKET NUMBER: 12
TELECOMMUNICATION INFORMATION:
TELEPHONE: 312/876-1800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       INFORMATION FOR SEQ ID NO: 14: SEQUENCE CHARACTERISTICS:
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Best Local Similarity 100..
2; Conservative
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TYPE: amino acid
STRANDEDNESS: single
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MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FRAGMENT TYPE: JORIGINAL SOURCE:
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USA
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         US-08-530-342A-14
                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
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NAME/KEY: AMIDATION
LOCATION: 5
OTHER INFORMATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for
                                                                  Patent No. 6127339
OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid
FORTER INFORMATION: synthesizer (9050 plus peptide synthesizer)
US-08-981-122-51
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APPLICANT: Bolognesi, Dani P.
APPLICANT: Matthews, Thomas J.
APPLICANT: Wild, Carl T.
APPLICANT: Wild, Carl T.
ITLE OF INVENTION: SYNTHETIC PEPTIDE INHIBITORS OF HIV TITLE OF INVENTION: TRANSMISSION
NUMBER OF SEQUENCES: 74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    65.2%; Score 15; DB 4; Length 5; ilarity 100.0%; Pred. No. 2e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                 Length 5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/554,616
FILING DATE: 06-NOV-1995
CLASSIFICATION: 435
PRICK APPLICATION NUMBER: US 08/073,028
FILING DATE: 07-JUN-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                           Query Match 65.2%; Score 15; DB 3;
Best Local Similarity 100.0%; Pred. No. 2e+05;
Matches 2; Conservative 0; Mismatches 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
STREET: 1155 Avenue of the Americas
CITX: New York
STATE: New York
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                          RESULT 20
US-08-554-616-44
; Sequence 44, Application US/08554616
; Patent No. 6133418
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Coruzzi, Laura A. 742
REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 78
TELECOMMUNICATION INFORMATION:
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INFORMATION POR SEQ ID NO: 44:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide
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Matches 2; Conserva
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Sequence 9, Application US/09208277;
Patent No. 6166177
GENERAL INFORMATION:
APPLICANT: Probst, Peter
APPLICANT: Skeiky, Yasir A.W.
TITLE OF INVENTION: COMPOUNDS AND METHODS FOR THE TREATMENT AND DIAGNOSIS
Sequence 14, Application US/08530342A
Patent No. 6143275
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: HIGER, CHRISTOPH-STEPH
APPLICANT: SCHIER, HANS-MARTIN
TITLE OF INVENTION: ACTIVE ISOTOPES, THEIR METAL COMPLEXES
TITLE OF INVENTION: TYPE S3N2 CHELATORS FOR RADIO
TITLE OF INVENTION: ACTIVE ISOTOPES, THEIR METAL COMPLEXES
NUMBER OF INVENTION: THEIR DIAGNOSTIC AND THERAPEUTICAL USE
NUMBER OF SEQUENCES: 15
CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: WOOD, PHILLIPS, VAN SANTEN, CLARK & ADDRESSEE: MORTIMER
                                                                                                                                                                                                                                                                                                                                                                                                              STREET: 500 West Madison Street, Suite 3800 CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE DISKETTE
COMPUTER: COMPAQ ProLinea 4/25s
COMPUTER: COMPAQ ProLinea 4/25s
SOFTWARE: WordPerfect for Windows 6.0a
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/530,342A
FILING DATE: 08-MAR-1996
CLASSIFICATION: 424
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/DE94/00370
FILING DATE: 29-MAR-1994
APPLICATION NUMBER: P 43 11 022.3
FILING DATE: 31-MAR-1993
ATTORNEY/AGENT INPRAMITON:
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APPLICANT: Barbas, Carlos
APPLICANT: Lerner, Richard A.
TITLE OF INVENTION: HETERODIMERIC RECEPTOR LIBRARIES USING
TITLE OF INVENTION: PHYSEMIDS
                                                                                                                                                                                                                                                                               Ouery Match 65.2%; Score 15; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 0; Mismatches 0; Indels
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CORRESPONDENCE ADDRESS:
ADDRESSEE: The Scripps Research Institute, Office of
ADDRESSEE: Patent Counsel
STREET: 10666 No. 6235469th Torrey Pines Road, TPC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
TITLE OF INVENTION: OF CHLAMYDIAL INFECTION FILE REFERENCE: 210121.469
CURRENT APPLICATION NUMBER: US/09/208,277
CURRENT FILING DATE: 1998-12-08
NUMBER OF SEQ ID NOS: 14
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 9
LENGTH: 5
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APPLICATION NUMBER: US 07/826,623
FILING DATE: 27-JAN-1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US 92/03091
FILING DATE: 10-APR-1992
ATTORNEY/AGENT INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/08/907,739
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; Sequence 148, Application US/08907739; Patent No. 6235469
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APPLICATION NUMBER: 08/133,011
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REFERENCE/DOCKET NUMBER: SCR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-554-2937
TELEPRAK: 619-554-6312
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SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
                                                                                                                                                                                         TYPE: PRT ORGANISM: Chlamydia trachomatis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: protein FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CA
USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: US
ZIP: 92037
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APPLICANT: Guthrie, K.
APPLICANT: Guthrie, K.
APPLICANT: Guthrie, K.
APPLICANT: Menutka, G.
APPLICANT: Menutka, G.
APPLICANT: Lambert, D.
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED PHARMACOKINETIC
TITLE OF INVENTION: HYBRE: US/09/315,304B
CURRENT FILING DATE: 1999-05-20
PRIOR PLIAGO DATE: 1999-05-20
PRIOR PLIAGO DATE: 1999-05-20
NUMBER OF SEQ ID NOS: 1667
SOFTWARE: PASCESEQ for Windows Version 3.0
SOFTWARE: PASCESEQ for Windows Version 3.0
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Ouery Match 65.2%; Score 15; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Anwer, Mohmed APPLICANT: Lambert, Dennis TITLE OF INVENTION: HYBRID POLYPEPTIDES WITH ENHANCED TITLE OF INVENTION: PHARMACOKINETIC PROFERTIES FILE REFERENCE: 7872-043
CURRENT APPLICATION NUMBER: US/09/082,279B
CURRENT FILING DATE: 1998-05-20
NUMBER OF SEQ ID NOS: 1515
SOFTWARE: FastSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 791, Application US/09315304B
; Patent No. 6348568
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                           ; Sequence 791, Application US/09082279B
; Patent No. 6288782
: GENEAL INFORMATION:
; APPLICANT: Barney, Shawn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTHER INFORMATION: Core polypeptide US-09-082-279B-791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Barney, Shawn
APPLICANT: Guthrie, Kelly
APPLICANT: Merutka, Gene
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Best Local Similarity
2, Conserva
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US-09-315-304B-791
                                                                                                                                                                                                                                RESULT 24
US-09-082-279B-791
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER SOFTWARE: FAST SEQ ID NO 791
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TELEFAX: 708-550...
INFORMATION FOR SEQ ID NO: 27
SEQUENCE CHARACTERISTICS:
THORME: 5 AMING acids
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CORRESPONDENCE ADDRESS:
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ZIP: 60064-3500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION:
   Patent No. 6448234
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                               4 LW 5
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STATE:
                                                                                                                                                                             SEQ ID NO 9
LENGTH: 5
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APPLICANT: Skeiky, Yaair
APPLICANT: Skeiky, Yaair
APPLICANT: Fling, Stewey Jeff
APPLICANT: Maisonneuve, Jeff
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR TREATMENT AND TITLE OF INVENTION: DIAGNOSIS OF CHLAMYDIAL INFECTION FILE REPREBUCE: 20121.469C5
CURRENT APPLICATION NUMBER: US/09/556,877
CURRENT PILING DATE: 2000-04-19
NUMBER OF SEQ ID NOS: 305
SOFTWARE: FastSEQ for Windows Version 3.0/4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       JOSEPS 298-594A-9

JESOPOLE 288-594A-9

Sequence 9, Application US/09288594A

Patent No. 644779

GENERAL INFORMATION:
APPLICANT: Probet, Peter
APPLICANT: Skelky, Yasir A.W.
APPLICANT: Skelky, Yasir A.W.
APPLICANT: Fling, Steeven P.
TITLE OF INVENTION: COMPOUNDS OF CHLAMYDIAL INFECTION
FILE REFERENCE: 210121.469C1
CURRENT APPLICATON NUMBER: US/09/288,594A

CURRENT FILING DATE: 1999-04-08

NUMBER OF SEQ ID NOS: 49

SOFTWARE: FastSEQ for Windows Version 3.0

LENGTH: 5
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 0; Indels
 0; Mismatches
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                                                                                                                                                        Sequence 9, Application US/09556877
Patent No. 6432916
GENERAL INFORMATION:
APPLICANT: Probst, Peter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-09-288-594A-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TYPE: PRT
; ORGANISM: Chlamydia trachomatis
US-09-556-877-9
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Conservative
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Best Local Similarity
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Best Local Similarity
Matches 2; Conserv
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US-09-620-412C-9
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US-09-556-877-9
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Matches
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APPLICANT: Severe P. FILING
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THEATMENT AND
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THEATMENT AND
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THEATMENT AND
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THEATMENT AND
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THEATMENT AND TYPE: PRI
COMPOSITION OF PRESENCE OF WINDOWS 195.000-07-20
COMPOSITION OF PART AND METHODS FOR THEIR USE
APPLICANT: TANKS C EXPRESS MEMBERS 195.000-07-20
COMPOSITION OF INVENTION OF MEAN AND METHODS FOR THEIR USE
CONTRY: MASSOCIATION DATE: 11. COUNTRY: USA
CONTRY: WHERE PRESENCE PRESS METHODS FOR THEIR USA
CONTRY: WHERE PRESS FOR METHODS FOR THEIR USA
CONTRY: WHERE PRESS FOR METHODS FOR THEIR USA
CONTRY: WHERE SHEED TO MENTOR THE STORY APPLICATION NUMBER: US/08/424.550
APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Patent Counsel
STREET: 10666 No. 6468738th Torrey Pines Road, TPC-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 161
CORRESPONDENCE ADDRESS:
ADDRESSEE: The Scripps Research Institute, Office of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMFUTER: IBM PC compatible
COMFUTER: IBM PC compatible
COMFUTER: IBM PC compatible
COMFUTER: IBM PC compatible
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
RAPPLICATION NUMBER: US/09/729,597
FILING DATE: 04-Dec-2000
CLASSIFICATION NUMBER: 08/133,011
APPLICATION NUMBER: 08/133,011
APPLICATION NUMBER: US 07/826,623
APPLICATION NUMBER: US 07/826,623
APPLICATION NUMBER: US 07/826,623
ATTORNEY/AGENT INFORMATION:
NAME: Fitting, Thomas
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: 34,163
REGISTRATION NUMBER: SCRF 238.2
TELEBPHONE: 619-554-2937
                                                                               Query Match 65.2%; Score 15; DB 4; Length 5; Best Local Similarity 100.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      65.2%; Score 15; DB 4; Length 5; 100.0%; Pred. No. 2e+05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid

TOPOLOGY: linear

MOLECULE TYPE: protein
FRAGMENT TYPE: internal
SEQUENCE DESCRIPTION: SEQ ID NO: 148:
                                                                                                                                                                                                                                                                                                   Sequence 148, Application US/09729597
Patent No. 6468738
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 148:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Kang, Angray
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
; TOPOLOGY: linear; MOLECULE TYPE: protein US-08-469-260A-228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             USA
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Best Local Similarity
Matches 2; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                                                                                               RESULT 30
US-09-729-597-148
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Gaps
                                                                                                      TITLE OF INVENTION: Altered Linolenic and Linoleic Acid Content
TITLE OF INVENTION: in Plants
NUMBER OF SEQUENCES: 72
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ·.
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APPLICANT: Takehana, Yoshi
APPLICANT: Ehresmann, Glenn
TITLE OF INVENTION: THERAPBUTIC STRATEGIES FOR
TITLE OF INVENTION: IMMUNOINFECTIVE CLUSTER VIRUS INFECTIONS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   65.2%; Score 15; DB 5; Length 5; ilarity 100.0%; Pred. No. 2e+05; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                     SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO) CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/01321
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CORRESPONDENCE ADDRESS:
ADDRESSEE: Robbins, Berliner & Carson
STREET: 201 North Figueroa Street, Suite 500
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                 JMBER: PCT/US94/01321
04-FEB-1994
                                                                                                                                                                                                                                                                                                                                         FILING DATE: 04-FEB-1994
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/156551
FILING DATE: 22-NOV-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/014431
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICATION NUMBER: US 08/029,850
FILING DATE: 11-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 103, Application PC/TUS9402631
; GENERAL INFORMATION:
RESULT 31
PCT-US94-01321-47
; Sequence 47, Application PC/TUS9401321
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          E: Floppy disk
IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                           COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Spitals, John P. REGISTRATION NUMBER: 29,215
                                                                                                                                                                                                        Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE: 05-FEB-1993
INFORMATION FOR SEQ ID NO: 4'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TOPOLOGY: linear;
; MOLECULE TYPE: peptide
PCT-US94-01321-47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 2; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                         Score 14; DB 1; Length 5;
Pred. No. 2e+05;
0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVENTION: RECOMBINANT THROMBIN RECEPTOR AND
TITLE OF INVENTION: RELATED PHARMACEUTICALS
NUMBER OF SEQUENCES: 223
CORRESPONDENCE ADDRESS:
ADDRESSE: MORRISON & FOERSTER
STREET: SOOO Pennsylvania Ave., NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORACTION NUMBER: US/08/475,263
PILING DATE: 07-JUN-1995
CLASSIFICATION: US/08/475,263
PILING DATE: 07-JUN-1995
CLASSIFICATION: US/08/475,263
PATTORNEY TINFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 22000-20502.03
TELECPHONE: (202) 887-1500
TELEPRX: (202) 887-0763
TELEFXX: 90-4030
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 83, Application US/08475263
; Patent No. 5759994
; GENERAL INFORMATION:
                   TELECOMMUNICATION INFORMATION:
                                                                                                                                               83:
                                          TELEPHONE: (415) 813-5600
TELEFAX: (415) 494-0792
TELEX: 34-0154
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                         Query Match 60.9%;
Best Local Similarity 40.0%;
Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: USA
ZIP: 2006-1812
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 40.0
Matches 2; Conservative
                                                                                                                                                                                                      LENGTH: 5 amino acids TYPE: AMINO ACID
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                                                                                                                                                                                                                             TYPE: AMINO ACLL
STRANDEDNESS: single
moboloGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: single
TOPOLOGY: linear
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US-07-789-184-83
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| Patent No. 5668768
| GENERAL INFORMATION:
| GENERAL INFORMATION:
| APPLICANT: SCARBOROUGH, ROBERT M. TITLE OF INVENTION: RECOMBINANT THROMBIN RECEPTOR AND TITLE OF INVENTION: RELATED PHARMACEUTICALS
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: ADDRESSEE
                                                                                                                                                                                                                                                                                                                                                                                  65.2%; Score 15; DB 5; Length 5; 100.0%; Pred. No. 2e+05; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    65.2%; Score 15; DB 6; Length 5; 100.0%; Pred. No. 2e+05; tive 0; Mismatches 0; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/789,184
FILING DATE: 19911107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 5171838

Patent No. 5171838

TITLE OF INVENTION: LEUJA BINDING PEPTIDES

NUMBER OF SEQUENCES: 24

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/526,921

FILING DATE: 22-MAY-1990

SEQ ID NO:3:

LENGTH: 5
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ATTORNEY/AGENT INFORMATION:
NAME: WIRASHIEE, KATE H.
REFERENCE/DOCKET NUMBER: 29.59
REFERENCE/DOCKET NUMBER: 22000-20502.20
REFERENCE/DOCKET NUMBER: 1920-331
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 977-1001
TELEPAX: (213) 977-1003
INFORMATION FOR SEQ ID NO: 103:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Conservative
                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: peptide
PCT-US94-02631-103
                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity Matches 2; Conserv
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Matches 2; Conserv
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US-07-789-184-83
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5171838-3
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                                                                                                                                                         APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVENTION: RELOADED PHARMACEUTICALS
TITLE OF INVENTION: RELATED PHARMACEUTICALS
NUMBER OF SEQUENCES: 223
CORRESPONDENCE ADDRESS:
ADDRESSE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVENTION: RECOMBINANT THROMBIN RECEPTOR AND
TITLE OF INVENTION: RELATED PHARMACEUTICALS
CORRESPONDENCE: 223
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/485,886

FILING DATE: 07-JUN-1995

CLASSIFICATION DATA:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 07/789,184

FILING DATE: 07-NOV-1991

ATTORNEY/AGENT INFORMATION:

NAME: REGISTRATION NUMBER: 22000-20502.20

TELERDANCE/DOCKET NUMBER: 22000-20502.20

TELERDANCE/DOCKET NUMBER: 22000-20502.20

TELERDANCE/DOCKET NUMBER: 22000-20502.20

TELERDANCE/DOCKET NUMBER: 24000

TELERDANCE/DOCKET NUMBER: 22000-20502.20

TELERDANCE/DOCKET NUMBER: 22000-20502.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                Sequence 83, Application US/08485886
Patent No. 5798248
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 83, Application US/08477362
Patent No. 5849507
GENERAL INFORMATION:
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STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
US-08-485-886-83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               94304-1018
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             94304-1018
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US-08-477-362-83
US-08-485-886-83
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 83, Application US/08477134
Patent No. 585648
GENERAL INFORMATION:
APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBOROUGH, ROBERT M.
APPLICANT: SCARBOROUGH, ROBERT M.
TITLE OF INVENTION: RECOMBINANT THROWBIN RECEPTOR AND
TITLE OF INVENTION: RELATED PHARMACEUTICALS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFEMILIAS 130-14. (2.05)/m3-05. (3.05) MB-1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: 05/08/477,134
FILING DATE: 07-01N-1995
CLASSIFICATION: 536
PRIOR APPLICATION: 536
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/789,184
FILING DATE: 07-NOV-1991
ATTORNEY/AGENT INFORMATION:
                    SOFTWARE: Patentin Release #1.0, Version #1.25
CUBRENT APPLICATION DAIR: US/08/477,362
FILICATION NUMBER: US/08/477,362
FILING DAIE: 07-JUN-1995
                                                                                                                                                                                                                                                                                               22000-20502.20
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                                                                                             FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/789,184
FILING DATE: 07-NOV-1991
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 223
CORRESPONDRAGE ADDRESS:
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ALIONALI MORASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REGISTRATION NUMBER: 22000
TELECOMMULICATION INFORMATION:
TELEPHONE: (415) 813-5600
TELERAX: (415) 494-0792
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                        NAME: MURSAHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFRENCE/DOCKET NUMBER: 2200C
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 494-0792
TELEX: 34-0154
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy o
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 755 ...
CITY: Palo Alto
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US-08-477-362-83
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GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 83, Application US/08473489A

Patent No. 6024936

GENERAL INFORMATION:
APPLICANT: COUGHLIN, SHAUN R.
TITLE OF INVENTION: RELATED PHARMACEUTICALS
ITLES OF INVENTION: RELATED PHARMACEUTICALS
ORRESPONDENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 755 Page Mill Road
CITY: Palo Alto
STATE: California
COUNTY: USA
                                                                                                                                                                       60.9%; Score 14; DB 2; Length 5;
40.0%; Pred. No. 2e+05;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 60.9%; Score 14; DB 3; Length 5; Best Local Similarity 40.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: THM PC compatible
COMPUTER: THM PC compatible
COMPUTER: THM PC compatible
COMPUTER: THM SAFEM: PC-DOS/MS-DOS
SOFTWARE: PRECATION DATA:
APPLICATION NUMBER: US/08/473,489A
FILING DATE: US/07/789,184
FILING DATE: US/07/789,184
FILING DATE: 1991-11-07
ATTORNEY/AGENT INFORMATION:
NAME: MURABAHIGE: KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 22000-20502.20
TELECOMMUNICATION INFORMATION:
MUST EDULATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 40
US-08-485-695-83
; Sequence 83, Application US/08485695
; Patent No. 6124101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (415) 494-0792
TELEX: 34-0154
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: (415) 813-5600
TELEFAX: (415) 494-0792
TYPE: amino acids TYPE: amino acids STRANDEDNESS: single COPPOLOGY: linear US-08-477-134-83
                                                                                                                                                 Query Match
Best Local Similarity 40.07
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: sin
TOPOLOGY: linear
                                                                                                                                                                                                                                                                1 FXXLW 5
                                                                                                                                                                                                                                                                                                           1 FEPFW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-473-489A-83
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Gaps
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| Patent No. 6197541
| GENERAL INFORMATION:
| GENERAL INFORMATION:
| APPLICANT: COUGHLIN, SHAUN R.
| TITLE OF INVENTION: RECOMBINANT THROMBIN RECEPTOR AND ITILE OF INVENTION: RELATED PHARMACEUTICALS
| VONBER OF SEQUENCES: 223
| CORRESPONDENCE ADDRESS:
| ADDRESSE: MORRISON & FOERSTER STREET: 755 Page Mill Road
| CITY: Palo Alto |
| CITY: Palo Alto |
| STATE: California |
| COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 60.9%; Score 14; DB 3; Length 5; Best Local Similarity 40.0%; Pred. No. 2e+05; Matches 2; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3; Indels
APPLICANT: COUGHLIN, SHAUN R.
APPLICANT: SCARBORGUGH, ROBERT M.
TITLE OF INVENTION: RECAMBINANT THROMBIN RECEPTOR AND
TITLE OF INVENTION: RELATED PHARMACEUTICALS
NUMBER OF SEQUENCES: 223
CORRESPONDENCE ADDRESS:
ADDRESSEE MORRISON & FOERSTER
STREET: 755 Page Mill Road
                                                                                                                                                                                                                                                                                                                                                                                OPERATING SYSTEM: PC-LUCS/PS-LUCS
SOFTWARE: Patentl NRelease #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/485,695
FILING DATE: 07-JUN-1995
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/789,184
FILING DATE: 07-NOV-1991
ATTORNEY/AGENT INPORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 22000-20502.20
TELEFONE: (415) 813-5600
TELEFONE: (415) 494-0792
TELEFAX: (415) 494-0792
TELEFAX: (415) 494-0792
TELEFAX: (5000-1000)
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 94304-1018
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      single
                                                                                                                                                                         STREET: 755 Page N
CITY: Palo Alto
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 FXXLW 5
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US-08-485-695-83
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4 LW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Matches
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APPLICANT: Yoshikawa, Kelichi
APPLICANT: Yoshowa, Kelichi
APPLICANT: Hassagawa, Masayasu
APPLICANT: Fujita, Hiroyuki
APPLICANT: Fujita, Hiroyuki
APPLICANT: Fujita, Hiroyuki
APPLICANT: Fujita, 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           60.9%; Score 14; DB 4; Length 5; 40.0%; Pred. No. 2e+05; tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/WS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/858,842
FILING DATE: 19920327
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: UP 142283
FILING DATE: 29-MAR-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: UP 298060
FILING DATE: 17-OCT-1991
ATYONEY/AGENT INFORMATION:
                                                                                                                                                         APPLICATION NUMBER: US 07/789,184
FILING DATE: 07-NOV-1991
ATTORNEY/AGENT INFORMATION:
NAME: MUTASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REPRENCE/DOCKET NUMBER: 22000-20502.20
TELEPHONE: (415) 813-5600
TELEPAN: (415) 813-5600
TELEFAX: 34-0154
INFORMATION FOR SEQ ID NO: 83:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: ARMSTRONG & KUBOVCIK
STREET: 1725 K Street N.W. Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Wasington
STATE: District of Columbia
COUNTRY: United States of America
APPLICATION NUMBER: US/08/018,760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               920247
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Patent No. 5314807
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REGISTRATION NUMBER: 34, 409
REFERENCE/DOCKET NUMBER: 920
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 659-2930
TELEFAX: (202) 887-0357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                 17-FEB-1993
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Matches 2; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                         PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
US-08-018-760-83
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                                             FILING DATE: 17
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 440142
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Yoshikawa, Masaaki
APPLICANT: Yoshikawa, Riichi
APPLICANT: Yoshikawa, Riichi
APPLICANT: Yoshikawa, Rouichi
APPLICANT: Hasegawa, Rouichi
APPLICANT: Yasumoco, Ryouichi
APPLICANT: Fujita, Hiroyuki
TITLE OF INVENTION: Method for Producing an Angiotensin
TITLE OF INVENTION: Converting Enzyme Inhibitor-Containing Composition
VOMBER OF SEQUENCES:
ADDRESSEE: Armstrong, Westerman, Hattori, McLeland &
ADDRESSEE: Armstrong, Westerman, Hattori, McLeland &
STREET: 1725 K Street, N.W., Suite 1000
                                                                                                                                                                                                                                                               Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0;
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                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                            56.5%; Score 13; DB 1; Length 5; 50.0%; Pred. No. 2e+05; cive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/960,636B
FILLING DATE: 14-OCT-1992
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: JP 298061/1991
FILLING DATE: 17-OCT-1991
ATTORNEY/AGENT INFRMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Tockman, Albert
REGISTRATION NUMBER: 19722
REFERENCE/DOCKET NUMBER: P1161-2679-A920798
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Mammalian meat, fish, crustaceans
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 2, Application US/07960636B
; Patent No. 5369015
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (202) 659-2930
TELEFAX: (202) 887-0357
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
INFORMATION FOR SEQ ID NO: 3:
                 SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: AMINO ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                               1; Conservative
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                                                                                        TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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Best Local Similarity
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                                                                                                                                                                    US-07-858-842-3
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                                                                                                                                                                                                                                                                                                            4 LW 5
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TOPOLOGY: UNKNOWN
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Best Local Similarity
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                                                                             NEW YORK
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                   ZIP: 10016
                                                          CITY: NEF
STATE: NI
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-07-802-667-19
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LOCATION: 5
OTHER INFORMATION: /label= Val-resin
OTHER INFORMATION: /note= "The carboxy terminal amino acid, Val, is
OTHER INFORMATION: attached to a resin"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           56.5%; Score 13; DB 1; Length 5; 50.0%; Pred. No. 2e+05; ative 1; Mismatches 0; Indels
                                                GENERAL INFORMATION:
APPLICANT: Heavner, George A.
APPLICANT: Rickinger, Douglas
APPLICANT: Mervic, Miljenko
TITLE OF INVENTION: Peptide Inhibitors of Selectin
TITLE OF INVENTION: Binding
NUMBER OF SEQUENCES: 13
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz &
ADDRESSEE: No. 5464935ris
STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                             COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compartible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/384,680
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CCOR-0024
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIPCATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 891,986
FILING DATE: 05-MAY-1992
ATTORNEY/AGENT INFORMATION:
NAME: Elderkin, Diame B.
REGISTRATION NUMBER: 28,588
REFERENCE/DOCKET NUMBER: CCOR-002
TELECOMONICATION: NIPORMATION:
TELECOMONICATION: 10-060-1304
TELEFAX: 215-568-3439
TELERX: 710-670-1334
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
                Sequence 8, Application US/08384680 Patent No. 5464935
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-133-020-4; Sequence 4, Application US/08133020; Patent No. 5476840; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: PIERRE BROTO, ET AL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: peptide
HYPOTHETICAL: NO
FEATURE:
                                                                                                                                                                                                                                                                                          STATE: Pennsylvania
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                 19103
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US-08-384-680-8
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1 IW 2
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TITLE OF INVENTION: NEW HEXPERTITOR.

TITLE OF INVENTION: DESIGNATION, METHOD AND TITLE OF INVENTION: DESIGNATIONS. SEREMATION, METHOD AND TITLE OF INVENTION: DESIGNATION: DESIGNATION SEREMATION: DESIGNATION: DESI
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0; Gaps
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APPLICANT: BEHANDY, PAUL W.
APPLICANT: LAMPE, JOHN W.
APPLICANT: MOHAMP, RAJU
APPLICANT: SHAW, KENNETH J.
TITLE OF INVENTION: INHIBITORS OF THE CONVERSION OF BIG
TITLE OF INVENTION: INHIBITORS OF THE CONVERSION OF BIG
TITLE OF INVENTION: BUDDTHELIN TO ENDOTHELIN
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: BERLEX LABORATORIES, INC.
STREET: 3300 FAIRFIELD ROAD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: 5
; COTHER INFORMATION: /note= "C-TERMINAL AMIDE ESTER"
US-07-802-667-19
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/802,667
FILING DATE: 19911205
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: BELLAMY, ELIZABETH A.
REGISTRACTION NUMBER: 27,854
REGISTRACTION NUMBER: BELLEX 51
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 10 PORMATION:
TELECOMMUNICATION 10 PORMATION 10 PORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: /note= "N-TERMINAL N-ACETYL'
                                                                                                                                                                                                                                            BERLEX 51
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                                         CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: BELLAMY, ELIZABETH A.
REGISTRATION NUMBER: 27,854
REFERENCE/DOCKET NUMBER: BERLE
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-305-472
INFORMATION FOR SEQ ID NO: 550
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 07470-5472
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5 amino acids
TYPE: AMINO ACID
STRANDEDNESS: single
FILING DATE: 19911205
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: peptide
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TITLE OF INVENTION: ENDOTHELIN TO ENDOTHELIN
NUMBER OF SEQUENCES: 33
CORRESPONDENCE ADDRESS:
ADDRESSEE: BERLEX LABORATORIES, INC.
                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                          56.5%; Score 13; DB 1; Length 5; 50.0%; Pred. No. 2e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                         ; OTHER INFORMATION: /note= "C-TERMINAL AMIDE ESTER"
US-07-802-667-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 07470-5472
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 18 PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/07/802,667
FILING DATE: 19911205
CLASSIFICATION: 514
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OTHER INFORMATION: /note= "N-TERMINAL N-ACETYL"
                                                                                                                                               LOCATION: 1
OTHER INFORMATION: /note= "N-TERMINAL N-ACETYL"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: BELLAMY, ELIZABETH A.
REGISTRATION NUMBER: 27,854
REPERENCE/POCKET NUMBER: BERLEX 51
TELECOMMUNICATION INFORMATION:
TELEPHONE: 201-305-5472
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 21, Application US/07802667; Patent No. 5504070
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: BIHOVSKY, RON H.
APPLICANT: ERHARDT, PAUL W.
APPLICANT: LAMPE, JOHN W.
APPLICANT: SHAW, KENNETH J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: Modified-site
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                                                                                                                         NAME/KEY: Modified-site
                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 50.08
Matches 1; Conservative
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LENGIH: 5 amino acids
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                       MOLECULE TYPE: peptide FEATURE:
                 TYPE: AMINO ACID
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide
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                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: USA
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                                                                                                                                                                                                                                LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-07-802-667-21
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                                                                                   Query Match 56.5%; Score 13; DB 1; Length 5; Best Local Similarity 50.0%; Pred. No. 2e+05; Matches 1; Conservative 1; Mismatches 0; Indels
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; LOCATION: 5
; OTHER INFORMATION: /note= "C-TERMINAL AMIDE ESTER"
US-07-802-667-21
                                                                                                                                                                                                                                                                     RESULT 49
US-08-057-167-12
i Sequence 12, Application US/08057167
j Patent No. 5541095
i GENERAL INFORMATION:
APPLICANT: Hischberg, Carlos B. APPLICANT: Hashimoto, Yasuhiro
APPLICANT: Swiedler, Stuart J. APPLICANT: Swiedler, Stuart J. APPLICANT: Swiedler, Stuart J. APPLICANT: Ishihara, Masayuki
TITLE OF INVENTION: GLYCOSAMINOGLYCAN SPECIFIC
ITLE OF INVENTION: SULFOTRANSFERASES
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: Massachusetts
CONNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER: U.S.A.
ZIP: 02110-2804
COMPUTER: SEADABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
COMPUTER: TBM PS/2 Model 502 or 555X
OPBEATING SYSTEM: MS-DOS (Version 5.0)
SOFTWARE: WordPerfect (Version 5.1)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/057,167
FILING DATE: 19930430
CLASSIFICIATION DATA:
APPLICATION NUMBER: 07/899,432
FILING DATE: 16 June 1992
ATTORNEY/AGENT INFORMATION:
NAME: CLARK, PAUL T.
REGISTRALION NUMBER: 30,162
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 04020/015002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617) 542-5070
TELEFAX: (617) 542-8906
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E: Fish & Richardson
225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 200154
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Conservative
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US-08-057-167-12
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STREET: 22
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RESULT 50 US-08-269-257-9

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Gaps
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US-08-373-911-5
Sequence 5, Application US/08373911
Patent No. 5663296
GENERAL INFORMATION:
APPLICANT: Doherty, Annette M.
TAPLICANT: Kornberg, Brian E.
APPLICANT: Nikam, Sham S.
TITLE OF INVENTION: Hydroxamate Inhibitors of
TITLE OF INVENTION: Endothecin Converting Enzyme
NUMBER OF SEQUENCES: 16
CORRESPONDENCES. 16
CORRESPONDENCE ADDRESS:
                Parent No. 5550110
GENERAL INFORMATION:
APPLICANT: Cody, Wayne L.
APPLICANT: Topliss, John G.
TILE OF INVENTION: Endothelin Antagonists II
NUMBER OF SEQUENCES: 64
CORRESPONDENCE ADDRESS:
ADDRESSEB: Warner-Lambert Company
STREET: 2800 Plymouth Rd.
                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Ver. #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/269,257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION 514
PROCR APPLICATION DATA:
PRIOR APPLICATION DATA:
PPLICATION NUMBER: US 08/033,515
FILING DATE: 31-MAR-1993
ATTORNEY/AGENT INFORMATION:
NAME: Tinney, Francis J.
REGISTRATION NUMBER: 33069
REFERENCE/DOCKET NUMBER: PD-4334-P2
TELECOMMUNICATION INFORMATION:
SUBJUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Warner-Lambert Company
                                                                                                                                                                                                                                                                                                                   ZIP: 48105
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
Sequence 9, Application US/08269257
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: sin
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Doherty, Annette M.
APPLICANT: Nornberg, Brian E.
APPLICANT: Nikam, Sham S.
TITLE OF INVENTION: Hydroxamate Inhibitors of
TITLE OF INVENTION: Endothecin Converting Enzyme
NUMBER OF SEQUENCES: 16
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/373,911
                  SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/373,911
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Warner-Lambert Company
STREET: 2800 Plymouth Rd.
CITY: Ann Arbor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/070,161
FILING DATE:
                                                                                   FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/070,161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Tinney, Francis J.
REGISTRATION NUMBER: 33069
REGISTRATION NUMBER: PD-4435
TELECOMMUNICATION INFORMATION:
TELEPHONE: 313 996-7295
                                                                                                                                                                                                                                                                   PD-4435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
WIDIUM TYPE: Floppy disk
CMBUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 52
US-08-373-911-6
; Sequence 6, Application US/08373911
; Patent No. 5663296
                                                                                                                                                                        FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Tinney, Francis J.
REGISTRATION NUMBER: 33069
REPERENCE/OCKET NUMBER: PD-4-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 313 996-7295
INFORMATION ESSO ID NO: 5:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 313 996-1553
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
TYPE: amino acid
TOPE: amino acid
TOPOLOGY: 1:--
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4 IW 5
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TITLE OF INVENTION: METHOD FOR DISCOVERY OF PEPTIDE AGONISTS
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: COOLEY GODWARD LLP
STREET: FIVE PALO ALTO SQUARE
STREET: FALO ALTO SQUARE
CITY: PALO ALTO STREET
STATE: CALIFORNIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           METHOD FOR DISCOVERY OF PEPTIDE AGONISTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             56.5%; Score 13; DB 2; Length 5; liarity 50.0%; Pred. No. 2e+05; Conservative 1; Mismarches
                                                                                                                                  0; Indels
                                                                                         56.5%; Score 13; DB 1; Length 5; 50.0%; Pred. No. 2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patent In Release #1.0, Version #1.25 CURRENT APPLICATION DATA: PAPLICATION DATA: PAPLICATION NUMBER: US/08/483,506A FILING DATE: 7-UNM-1995 CLASSIFICATION: 435
                                                                                                                                1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           UCAL-239/00US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 2, Application US/08483506A
Patent No. 5925529
GENERAL INFORMATION:
APPLICANT: COUGHLIN, ET AL.
ITLE OF INVENTION: METHOD FOR DI
                                                                                                                                                                                                                                                                                                           Sequence 1, Application US/08483506A
patent No. 5925529
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: COOLEY GODWARD LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ATTORNEY/AGENT INFORMATION:
NAME: NEELEY, RICHARD L. PhD
REGISTATION NUMBER: 30,092
REFERENCE/DOCKET NUMBER: UCAL
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 843.5000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: (415) 857-0663
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                          Query Match
Best Local Similarity 50.0%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
            ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-373-911-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 1. CORRESPONDENCE ADDRESS:
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Matches 1; Conserv
TYPE: amino acid
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                                                                                                                                                                                                                                                                       RESULT 53
US-08-483-506A-1
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                                                                                                                                                                                               : 4
1W 5
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Matches
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Patent No. 595579

GENERAL INFORMATION:
APPLICANT: Leonard, James N. Montminy, Marc R.
TITLE OF INVENTION: REGULATOR OF INSULIN GENE EXPRESSION
TITLE OF INVENTION: REGULATOR OF INSULIN GENE EXPRESSION
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSE: Klauber & Jackson
STREET: 411 Hackensack Avenue
CITY: Hackensack
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                         COUNTRY: USAS
ZIP: 94306-2185
ZIP: 94306-2185
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patent In Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/483,506A
FILING DATE: 7-JUN-1995
ATTORNET/AGENT INFORMATION:
NAME: NEELEY, RICHARD L. PhD
REGISTRATION NUMBER: 30,092
REFERENCE/DOCKET NUMBER: UCAL-239/00US
FELECOMMONICATION INFORMATION:
TELEPHONE: (415) 843-5000
TELECOMMONICATION INFORMATION:
TELEPHONE: (415) 843-5000
TELEFAX: (415) 843-5000
TELEFAX: (415) 87-0663
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
FENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LOCATION: 1
OTHER INFORMATION: /note= "Should be Formyl-Met'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/437,607A
FILING DATE: MAX 9, 1995
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 1110-1-001 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
: FIVE PALO ALTO SQUARE
: 3000 EL CAMINO REAL
PALO ALTO
CALIFORNIA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: Modified-site
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Hackensack
STATE: New Jersey
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
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Matches 1; Conserv
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US-08-437-607A-38
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STREET:
STREET:
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; Patent No. 6127339
; OTHER INFORMATION: F-moc amino acids by solid phase method using a multipeptide
; OTHER INFORMATION: synthesizing system (RaMPS)
US-08-981-122-2
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OTHER INFORMATION: Sequence of a peptide synthesized in Example 1 from L-form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/08/981,122B
CURRENT FILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: P7-176904
PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR PLING DATE: 1996-06-21
PRIOR PLING DATE: 1996-06-21
SPRIOR FLING DATE: 1996-06-21
SPRIOR FLING DATE: 1996-06-21
SUMBER OF SEQ ID NOS: 90
SOFTWARE: PatentIn Ver: 2.0
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APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Hatanaka, Masaharu
ITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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0
                                                                                                                                              TYPE: amino acid
TOPOLOGY: Linear
MOLECULE TYPE: proctein
DESCRIPTION: amino acids 47-51 of the homeobox homeodomain
                                                                                                                                                                                                                                                                                                                                                       56.5%; Score 13; DB 2; Length 5; 50.0%; Pred. No. 2e+05;
                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                         1; Mismatches
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CURRENT FILING DATE: 1997-12-18
PRIOR APPLICATION NUMBER: JP 7-176904
PRIOR FILING DATE: 1995-06-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 19, Application US/08981122B
; Patent No. 6127339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 2, Application US/08981122B
; Patent No. 6127339
; GENERAL INFORMATION:
                     TELEFAX: 201 343-1684
TELEX: 133521
INFORMATION FOR SEQ ID NO: 38:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: PRT
ORGANISM: Artificial Sequence
  201 487-5800
                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                            5 amino acids
                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 1; Conserv
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                                                                                                                                                                                                                                                                                  ORGANISM: Rat
                                                                                                                                                                                                                                                        ORIGINAL SOURCE:
TELEPHONE:
                                                                                                                                                                                                                                                                                                     US-08-437-607A-38
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LENGTH: 5
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LOCATION: 5
OTHER INFORMATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for
                                                                                                                                                                                                                                                                                                                                                               Patent No. 6127339
OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid
OTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Sequence of a peptide synthesized in Comparative Example 2 from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Patent No. 6127339
OTHER INPORMATION: L-form F-moc amino acids by solid phase method using a CTHER INFORMATION: multipeptide synthesizing system (RAMPS)
US-08-981-122-72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
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CURRENT PILING DATE: 1997-12-18
CURRENT FILING DATE: 1995-06-21
PRIOR FILING DATE: 1995-06-21
PRIOR FILING DATE: 1996-06-21
PRIOR FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: PatentIn Ver: 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
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               PRIOR APPLICATION NUMBER: JP 7-176904
PRIOR FILING DATE: 1995-06-21
PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver. 2.0
LENGTH: 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , Sequence 72, Application US/08981122B
; Patent No. 6127339
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 4, Application US/08871600A
; Patent No. 6222021
; GENERAL INFORMATION:
CURRENT FILING DATE: 1997-12-18
                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                              NAME/KEY: AMIDATION
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US-08-981-122-72
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                                                                                                                                                                                                                           TYPE: PRT
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LOCATION: 5

LOCATION: 5

LOCATION: Sequence of peptides synthesized in Examples 11 and 12 from L-for patent No. 6127339

OTHER INFORMATION: F-moc amino acids by solid phase method using an automatic peptid OTHER INFORMATION: synthesizer (9050 plus peptide synthesizer)

US-08-981-122-37
                                                                                                                                                                                                                                           Patent No. 6127339
OTHER INFORMATION: F-moc amino acids by solid phase method using a multipeptide
OTHER INFORMATION: synthesizing system (RaMPS)
                                                                                                                                                                                                                         OTHER INFORMATION: Sequence of a peptide synthesized in Example 2 from L-form
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  .
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APPLICANT: Hatanaka, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
FILE REFERENCE:
CURRENT APPLICATION NUMBER: US/08/981,1228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Hatanaka, Yoshihiro
APPLICANT: Aritomi, Masaharu
TITLE OF INVENTION: Peptide for binding thereto a low density lipoprotein
                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                    56.5%; Score 13; DB 3; Length 5; 50.0%; Pred. No. 2e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/08/981,122B
CURRENT FILING DATE: 1997-12-18
CURRENT FILING DATE: 1995-06-21
PRIOR APPLICATION NUMBER: UP 7-176504
PRIOR FILING DATE: 1995-06-21
PRIOR FILING DATE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: Patentin Ver: 2.0
PRIOR APPLICATION NUMBER: PCT/JP96/01734
PRIOR FLING DAFE: 1996-06-21
NUMBER OF SEQ ID NOS: 90
SOFTWARE: PATENTIN VEY. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 37, Application US/08981122B; Patent No. 6127339; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 53, Application US/08981122B
; Patent No. 6127339
                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
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tes 1; Conserva
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                                                                                               SEQ ID NO 19
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                                                                                                                              LENGTH:
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4 LW 5
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Patent No. 6268328
GENERAL INFORMATION:
APPLICANT: Mitchinson, Colin
APPLICANT: Wendt, Dan J.
TITLE OF INVENTION: No. 6268328el Variant EGIII-Like Cellulase Compositions
FILE REPERBNCE: GC555
CURRENT APPLICATION NUMBER: US/09/216,295
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TITLE OF INVENTION: Endotoxin Binding and Neutralizing TITLE OF INVENTION: Protein and Uses Thereof NUMBER OF SEQUENCES: 4
CORRESPONDENCE ABESES: ADDRESSES: Sterne, Kessler, Goldstein & Fox P.L.L.C. STREET: 1100 New York Avenue, N.W., Suite 600
                                                                                                                  1413.001000A/RWE
                                                                               CITY: Washington
STATE: District of Columbia
COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Esmond, Robert W. REGISTRATION NUMBER: 32,893
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REFERENCE/DOCKET NUMBER: 14.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEFAX: (202) 371-2540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEX: 248636 SSK
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS: single
TOPOLOGY: not relevant
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Best Local Similarity 50.v.
1, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , MOLECULE TYPE: peptide US-08-871-600A-4
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TITLE OF INVENTION: ARTIFICIAL ANTIBODY POLYPEPTIDES
NUMBER OF SEQUENCES: 118
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, Ste. 1600
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                           56.5%; Score 13; DB 4; Length 5; ilarity 50.0%; Pred. No. 2e+05; Conservative 1; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQ ID NOS: 41
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 26
LENGTH: 5
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                                                                                                                                                            LOCATION: (1) ... (5)
OTHER INFORMATION: Xaa = Leu, Phe, or Ile
                                                                                                                                                                                                                                                                                                                                                                                                          ; Sequence 51, Application US/09638202A
; Patent No. 6462189
; GENERAL INFORMATION:
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CURRENT FILING DATE: 1998-12-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
                                                                                TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 55402
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 50.0
Matches 1; Conservative
                                                                                                                               OTHER INFORMATION: Peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        HYPOTHETICAL: NO ANTI-SENSE: NO
                                                                                                                                                                                                                                         Best Local Similarity
Matches 1; Conserv
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                                                                                                                                                NAME/KEY: VARIANT
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US-09-638-202A-51
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COUNTRY:
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                                                                                                                                                                    APPLICANT: Hirschberg, Carlos B.
APPLICANT: Orellana, Ariel
APPLICANT: Orellana, Ariel
APPLICANT: Swiedler, Yasuhiro
APPLICANT: Swiedler, Stuart J.
APPLICANT: Wei, Zheng
APPLICANT: Ishinara Masayuki
TITLE OF INVENTION: GLYCOSAMINOGLYCAN SPECIFIC
TITLE OF INVENTION: SULFOTRANSFERASES
NUMBER OF SEQUENCES: 19
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Cody, Wayne L.
APPLICANT: Doherty, Annette M.
APPLICANT: Topliss, John G.
TITLE OF INVENTION: Endothelin Antagonists II
CORRESPONDENCES: 64
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 07/899,432
FILING DATE: 16 June 1992
ATTORNEY/AGENT INFORMATION:
NAME: CLARK, PAUL T.
REGISTRATION NUMBER: 30,162
REFERENCE/DOCKET NUMBER: 04020/015002
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHONE: (617) 542-5070
TELEFRAX: (617) 542-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: 3.5" Diskette, 1.44 Mb COMPUTER: 1BM PS/2 Model 50Z or 55SX OPREATING SYSTEM: MS-DOS (Version 5.0) SOFTWARE: WordPerfect (Version 5.1) APPLICATION DATA: APPLICATION DATA: PCT/US93/05412 FILING DATE: 19930607
                                                                                                                        Sequence 12, Application PC/TUS9305412
GENERAL INFORMATION:
APPLICANT: Hirschberg, Carlos B.
APPLICANT: Orellana, Ariel
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GENERAL INFORMATION:
APPLICANT: Cody, Wayne L.
                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Fish & Richardson
STREET: 225 Franklin Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELERAX: (617) 542-8906
TELEX: 200154
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS;
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Best Local Similarity 50.0°
Matches 1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Dis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acid STRANDEDNESS: N/H
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                                                                                                                                                                                                                                                                                                                                                                                                               Boston
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                    2 MW 3
4 LW 5
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Best Local Similarity 50.0%; Pred. No. 2e+05;
Matches 1; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: KAUVAR, LAWRENCE M. TITLE OF INVENTION: METHOD TO PRODUCE IMMUNODIAGNOSTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5217869-22; Patent No. 5217869; APPLICANT: KAUVAR, LAWRENCE M.; TITLE OF INVENTION: METHOD TO PRODUCE IMMUNODIAGNOSTIC
                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Ver. #1.25
CURRENT APPLICATION DATA:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                    NAME: TINNEY, FRANCIS J.
REGISTRATION NUMBER: 33069
REFERENCE/DOCKET NUMBER: PD-4334-P2
TELECHONE: 313 996-7295
TELEPHONE: 313 996-7295
INPORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS:
ADDRESSEE: Warner-Lambert Company
STREET: 2800 Plymouth Rd.
CITY: Ann Arbor
                                                                                                                                                                                                                                                                                                                                         US 08/033,515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
                                                                                                                                                                                                                                                                                            CLASSIPICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 06
FILING DATE: 31-MAR-1993
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               5 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear MOLECULE TYPE: peptide PCT-US95-04171-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 121
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;Patent No. 5217869
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Brook, Smith & Reynolds, P.C.
                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMFUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
   ADDRESSEE: Hamilton, Broo.
STREET: Two Militia Drive
CITY: Lexington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: AMINO ACID
MOLECULE TYPE: peptide
                                                                                                 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
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                                                                   COUNTRY: US
ZIP: 02173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 71
US-08-290-448A-3
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                                                                                                                56.5%; Score 13; DB 6; Length 5; 50.0%; Pred. No. 2e+05; ative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     56.5%; Score 13; DB 6; Length 5; 50.0%; Pred. No. 2e+05; Live 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 56.5%; Score 13; DB 6; Length 5; Best Local Similarity 50.0%; Pred. No. 2e+05;
                                                                                                                                                                                                                                                                         5217869-24
;Patent No. 5217869
; APPLICANT: KAUVAR, LAWRENCE M.
; TITLE OF INVENTION: METHOD TO PRODUCE IMMUNODIAGNOSTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AE27869-34
; Patent No. 5217869
; APPLICANT: KAUVAR, LAWRENCE M.
; TITLE OF INVENTION: METHOD TO PRODUCE IMMUNODIAGNOSTIC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 4, Application US/07946237
Patent No. 5348874
GENERAL INFORMATION:
GENERAL INFORMATION:
SAPLICANT: Savakis, Charalambos
APPLICANT: Franz, Gerald H
APPLICANT: Loukeris, Athanasios
TITLE OF INVENTION: Eukaryotic Transposable Element
NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Mismatches
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
                                                                                                                                                                                                                                                                                                                                                                           CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/255,906
FILING DATE: 11-OCT-1988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Conservative
                                                                                                                                                Conservative
                                                                                               Query Match
Best Local Similarity
1; Conserva
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 5
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5217869-34
                                                                 LENGTH: 5
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1 MW 2
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                                                  ,SEQ ID NO:22:
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Sequence 3, Application US/08290448A

Patent No. 5676954

GENERAL INFORMATION:
APPLICANT: Raper, Bavid G.
APPLICANT: Rafner, Thorunn
APPLICANT: Raner, Mai-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
CORRESONDENCES: 93
CORRESONDENCES: 93
STREET: 60 State Street, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 5;
CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/07/946,237

FILING DATE: 19920914

CLASSIFICATION: 435

ATTONNEY/AGENT INPORMATION:

NAME: David E. Brock

REGISTRATION NUMBER: 22,592

REGISTRATION NUMBER: 22,592

REGISTRATION NUMBER: BTT92-01

TELECOMMUNICATION INPORMATION:

TELEPHONE: (617) 861-6240

TELEPHONE: (617) 861-6240

INFORMATION FOR SEQ ID NO: 4:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/290,448A
FILING DATE: August 15, 1994
PRIOR APPLICATION NUMBER: US/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/529,951
FILING DATE: March 17, 1989
ATTORNEY/AGENT INFORMATION:
NAME: Amy E Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IM-018CN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 52.2%; Score 12; DB 1;
Best Local Similarity 50.0%; Pred. No. 2e+05;
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Gaps

1 VW 2

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RESULT 73
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Patent No. 5676954
GENERAL INFORMATION:
APPLICANT: Rogers, Bruce
APPLICANT: Klapper, David G.
APPLICANT: Kafnar, Thorunn
APPLICANT: Kuo, Mel-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAHIVE & COCKFIELD
STREET: 60 State Street, suite 510
                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                  52.2%; Score 12; DB 1; Length 5; ilarity 50.0%; Pred. No. 2e+05; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                52.2%; Score 12; DB 1; Length 5; 50.0%; Pred. No. 2e+05; tive 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
PLICATION NUMBER: US/08/290,448A
FILING DATE: August 15, 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI-018CN
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
TELEPHONE: (617)227-5941
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: MArch 17, 1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                 TELEPHONE: (617)227-7400
TELEPRAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
TYPE: amino acid
TOPOLOGY: linear
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Massachusetts
COUNTRY: USA
ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 50...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                         MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
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Best Local Similarity
Matches 1; Conserv
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US-08-290-448A-39
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Gaps
                                                                                  APPLICANT: Kemp, Bruce E.
APPLICANT: Nicholson, Geoffrey C.
APPLICANT: Martin, Thomas J.
APPLICANT: Fenton, Anna J.
APPLICANT: Hammonds, R. Glenn
TITLE OF INVENTION: COMPOUNDS AND COMPOSITIONS WHICH INHIBIT
TITLE OF INVENTION: BONE RESORPTION
NUMBER OF SEQUENCES: 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          52.2%; Score 12; DB 1; Length 5; 20.0%; Pred. No. 2e+05; tive 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Indels
                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert, ADDRESSEE: Attn: W.H. Dreger
STREET: 4 Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/064,111C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/064,111C
FILING DATE: 12-AUG-1993
CLASSIPICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/AU91/00580
FILING DATE: 13-DEC-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU PK9567
FILING DATE: 19-NOV-1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: AU PK3879
FILING DATE: 13-DEC-1990
ATTORNEY/AGENT INFORMATION:
NAME: DYSECTION NUMBER: AU PK3879
FILING DATE: 13-DEC-1990
ATTORNEY/AGENT INFORMATION:
NAME: DYSECTION NUMBER: 24,190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      A-58456/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
            Sequence 22, Application US/08064111C; Patent No. 5688760
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 3, Application US/08290448A, Patent No. 5698204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: A-TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 22:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                         RY: United States 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 20.0 Matches 1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
                                                                     GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS:
US-08-064-111C-22
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Gaps

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0; Indels

4 LW 5

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Job time : 15 secs
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| Patent No. 5698204
| GENERAL INFORMATION:
| APPLICANT: RAGBER, David G.
| APPLICANT: RAIDPER, David G.
| TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
| TITLE OF SEQUENCES: 93
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: LAHIVE & COCKFIELD
| STREET: 60 State Street, suite 510
| CITY: Boston
| CATATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ô
                   APPLICANT: Rogers, Bruce
APPLICANT: Klapper, David G.
APPLICANT: Klapper, David G.
APPLICANT: Kuo, Mei-chang
TITLE OF INVENTION: Allergenic Proteins From Ragweed and Uses
NUMBER OF SEQUENCES: 93
CORRESPONDENCE ADDRESS:
ADDRESSER: LAHITE & COCKFIELD
STREET: 60 State Street, suite 510
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   52.2%; Score 12; DB 1; Length 5; 50.0%; Pred. No. 2e+05; ative 1; Mismatches 0; Indels
                                                                                                                                                                                                                         CITY: Massachusetts
COUNTRY: USA
ZIP: D2109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: INDA
COMPOTARE: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/290,448A
FILING DATE: August 15, 1994
FILING DATE: August 15, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/529,951
FILING DATE: May 29, 1990
APPLICATION NUMBER: US 07/325,365
FILING DATE: March 17, 1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Amy E. Mandragouras
REGISTRATION NUMBER: 36,207
REFERENCE/DOCKET NUMBER: IMI
TELECOMMUNICATION INFORMATION:
TELEPHONE: (617)227-7400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 02109-1875
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (617)227-5941
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 5 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-290-448A-39
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-290-448A-3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 VW 2
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/290,448A
FILNG DATE: August 15, 1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/529,951
FILING DATE: March 17, 1989
ATTORNEY/AGENT INFORMATION:
NAME: Amy E. Mandragourae
REGISTRATION NUMBER: 36,207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Search completed: February 12, 2003, 10:53:11
                                                                                                                                                                                                                                                                                      REFRENCE/DOCKET NUMBER: INI-018CN
TELECOMMUNICATION INFORMATION:
TELEPAK: (617)227-5941
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 5 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide
FRAGMENT TYPE: internal
                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-290-448A-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: ami
TOPOLOGY:
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GenCore version 5.1.3 (c) 1993 - 2003 Compugen Ltd.
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OM protein - protein search, using sw model

February 12, 2003, 11:12:22 ; Search time 5.41667 Seconds (without alignments) 88.739 Million cell updates/sec Run on:

US-09-403-440A-1 23 Title: Perfect score:

1 PXXLW 5 Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

283224 segs, 96134422 residues Searched:

4984 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 25

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

PIR 73:*
1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

STRAMATES

		de			SUMMARIES	
Result No.	Score	Query Match	/ n Length	DB	ID	
1	20		0 15	5	PH1613	
7	17		91 6	~	C37290	homeotic protein G
м	17	ë.	9 25	~	868897	phospholipase A2 (
4	16	•		~	S38281	GTP-binding protei
ഹ	15	ω.		~	B34835	dnaA protein - Pse
9	15	•		7	PT0519	T-cell receptor be
7	15	S.		~	PH1602	J-0-
ω	15	'n.		~	A39308	glycine reductase
თ	15	S.		7	A24244	adipokinetic hormo
10	15			7	S07241	litorin - Rohde's
11	15	ů.	2 10	~	E49033	T-cell receptor ga
12	15	'n.		7	F49033	T-cell receptor ga
13	15	65.3	2 11	Н	LFTWWE	probable trpEG lea
14	15	δ.		~	C53652	rhlR protein - Pse
15	15	•		~	A29169	phospholipase A2 (
16	15	ت	2 12	~	164829	gene HEXA protein
17	15	ů.		~	A26093	microbial collagen
18	15	δ.	2 12	~	158273	thyroglobulin - ra
19	15	'n.		~	PH1620	Ig H chain V-D-J r
50	15	65.5		~	G37266	Ig heavy chain C r
21	15	ď.		7	D37267	Ig heavy chain C r
22	15	S.		~	151905	collecting duct wa
23	15	Ŋ.		~	S54344	glyceraldehyde-3-p
24	15	S.		N	A35105	hypothetical prote
25	15	Ŋ.		N	PH1625	Ig H chain V-D-J r
56	15	65.2	2 14	~	PH1626	Ig H chain V-D-J r
27	15	رى س		~	PH1627	H chain
28	15	ů.		~	PH1594	Ig H chain V-D-J r
53	15	•	2 14	7	PH0801	[e]

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Gaps

.; 0

Query Match 73.9%; Score 17; DB 2; Length 16; Best Local Similarity 40.0%; Pred. No. 1e+03; Matches 2; Conservative 1; Mismatches 2; Indels

12 FKKMW 16

1 FXXLW 5

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ALIGNMENTS

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Ig H chain V-D-J region (clone B-less 17) - mouse (fragment)
(Species: Mus musculus (house mouse)
(Spacies: Mus musculus (house mouse)
(Spacies: 02-Jun-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
(SAccession: PH1613
(R;Levinson, D.A.; Campos-Torres, J.; Leder, P.
A;Levinson, D.A.; Campos-Torres, J.; Leder, P.
A;Title: Molecular characterization of transgene-induced immunodeficiency in B-less mice
A;Reference number: PH1580; MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cypecies: Mus musculus (house mouse)
Cypecies: Musculus (house)
Cypecies: Musculus (ho
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Molecule type: DNA
A;Residues: 1-15 cLEV>
A;Experimental source: bone marrow pre-B lymphocyte
C;Keywords: immunoglobulin
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Matches 3; Conserv
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CiSpecies: Mus musculus (house mouse)
CiSpecies: Mus musculus (house mouse)
CiDate: O2-Unn-1994 #sequence_revision 02-Jun-1994 #text_change 17-Mar-1999
CiAccession: PH1602
Rilevinson, D.A.; Campos-Torres, J.; Leder, P.
Rilevinson, D.A.; Campos-Torres, J.; Leder, P.
A;Title: Modecular characterization of transgene-induced immunodeficiency in B-less mice
A;Reference number: PH1580; MUID:93301609; PMID:8315387
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         glycine reductase (EC 1.4.99.-) sulfhydryl protein C, alpha chain - Clostridium sticklan
C,Species: Clostridium sticklandii
C,Date: 19-Jun-1992 #sequence_revision 19-Jun-1992 #text_change 15-Aug-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A,Description: glycine reductase complex catalyzes the reductive deamination of glycine C;Keywords: ATP; oxidoreductase
                                                                                                                                                                                                                                                                                                                     J. Exp. Med. 174, 115-124, 1991
A;Title: Junctional sequences of fetal T cell receptor beta chains have few N regions. A;Reference number: PT0509; MUID:91277601; PMID:1711558
A;Reference: PT0519
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                 T-cell receptor beta chain V-D-J region (100-4C) - mouse (fragment)
C;Species: Mus musculus (house mouse)
C;Date: 17-Jul-1992 #sequence_revision 17-Jui-1992 #text_change 30-May-1997
C;Accession: PT619
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 2; Length 6; 100.0%; Pred. No. 2.8e+05; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Experimental source: bone marrow pre-B lymphocyte C; Keywords: immunoglobulin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Experimental source: adult thymus, strain BALB/c
                                                                                                                                                                                                                                                                                                                                                                                                                                    A;Status: translation not shown A;Molecule type: mRNA
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A;Molecule type: protein
A;Residues: 1-8 <STA>
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Matches 2; Conserva
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A; Residues: 1-7 < LEV>
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                                                                                                               C;Accession: S68897; """" Campbell, D.G.; Cohen, P.; Creaney, A.; Wilton, D.C.; Ma Bur. J. Biochem. 238, 690-697, 1996
A;Title: Cloning and expression of cytosolic phospholipase A(2) (cPLA(2)) and a naturall A;Reference number: S68897; WUID:96300233; PMID:8706669
A;Accession: S68897
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Accession: B14835
R;Yee, T.W.; Smith, D.W.
Proc. Natl. Acad. Sci. U.S.A. 87, 1278-1282, 1990
A;Title: Pseudomonas chromosomal replication origins: a bacterial class distinct from Es A;Accession: B34835, MUID:90160310; PMID:2106132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      C; Accession: S38281
R; Moore, M.S.; Blobel, G.
Nature 365, 661-663, 1993
A; Title: The GTP-binding protein Ran/TC4 is required for protein import into the nucleus A; Reference number: S38281; MUID: 94019818; PMID: 8413630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: protein
A;Mesidues: 1-24 eMOO>
C;Superfamily: ras transforming protein; translation elongation factor Tu homology
C;Keywords: GTP binding
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                                                phospholipase A2 (BC 3.1.1.4), cytosolic - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 04-Dec_1997 #sequence_revision 12-Dec-1997 #text_change 17-Mar-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Xenopus sp. (clawed frog)
C;Date: 19-May-1994 #sequence revision 27-Feb-1997 #text_change 28-May-1999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            C,Species: Pseudomonas aeruginosa
C,Date: 13-Jul-1990 #sequence_revision 13-Jul-1990 #text_change 08-Oct-1999
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A;Residues: 1-6 <YEE>
A;Cross-references: GB:M30125; NID:g151419; PIDN:AAA25916.1; PID:g151421,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        73.9%; Score 17; DB 2; Length 25; 40.0%; Pred. No. 1.5e+03; Live 1; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                              A;Molecule type: mRNA
A;Residues: 1-25 <GOR>
C;Keywords: carboxylic ester hydrolase; cytosol
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GTP-binding protein - clawed frog (fragments)
                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity 40.0%
2, Conservative
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A; Status: preliminary
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Gaps

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probable trpEG leader peptide - Thermus aquaticus Cispecies: Thermus aquaticus Cispecies: Thermus aquaticus Cispecies: Thermus aquaticus Cipate: 30-Jun-1991 #sequence_revision 30-Jun-1991 #text_change 16-Jul-1999 CiAccession: 803315 Risato, S; Nakada, Y; Kanaya, S.; Tanaka, T. Risato, S; Nakada, Y.; Kanaya, S.; Tanaka, T. Aititle: Molecular Acta 950, 303-312, 1988 Aititle: Molecular cloning and nuclectide sequence of Thermus thermophilus HB8 trpE and A;Reference number: 803315; MUID:89000781; PMID:2844259
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          T-cell receptor gamma chain V-D-J region - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Species: 19-Dec-1993 #sequence_revision 17-Mar-2000 #text_change 17-Mar-2000
C;Accession: F49033
R;Morita, C.T.; Verma, S.; Aparicio, P.; Martinez, C.; Spits, H.; Brenner, M.B.
Eur. J. Immunol. 21, 2999-3007, 1991
A;Title: Functionally distinct subsets of human gamma/delta T cells.
A;Reference number: A49033; MUID:92083926; PMID:1684157
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A,Residues: 1-11 <SAT>
A,Cross-references: EMBL:X07744; NID:g48261; PIDN:CAA30565.1; PID:g48262
A,Note: the source is designated as Thermus thermophilus HB8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A;Cross-references: GB:S72605; NID:g240700; PIDN:AAB20632.1; PID:g240701
A;Note: sequence extracted from NCBI backbone (NCBIN:72605, NCBIP:72606)
C;Keywords: T-cell receptor
                                                                                                                                                                                                                                                                                                                    A,Molecule type: DNA
A;Residues: 1-10 <MOR2>
A;Cross-references: GB:S72587; NID:g240696; PIDN:AAB20630.1; PID:g240697
A;Note: sequence extracted from NCBI backbone (NCBIN:72587, NCBIP:72589)
C;Keywords: T-cell receptor
                                                                                                                                                                    A;Residues: 1-10 <MOR>
A;Residues: 1-10 <MOR>
A;Croses.references: GB:S72587; NID:g240696; PIDN:AAB20630.1; PID:g240697
A;Note: sequence extracted from NCBI backbone (NCBIN:72591, NCBIP:72595)
A;Accession: D49033
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0
A, Title: Functionally distinct subsets of human gamma/delta T cells. A, Reference number: A49033; MUID:92083926; PMID:1684157 A, Accession: E49033 A, Status: preliminary
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100.0%; Pred. No. 1.8e+03;
tive 0; Mismatches 0; Indels
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ilarity 100.0%; Pred. No. 1.8e+03;
Conservative 0; Mismatches 0;
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A; Molecule type: DNA
A; Residues: 1-10 < MOR>
                                                                                                                                                                                                                                                                                          A; Status: preliminary
                                                                                                                                       A, Molecule type: DNA
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Cîspecies: Heliothis zea (bollworm, corn earworm, tomato fruitworm)
Cjate: 31-Mar-1988 #sequence_revision 23-Mar-1995 #text_change 31-Oct-1997
CjAccession: A2444
RjJaffe, H.; Raina, A.K.; Riley, C.T.; Fraser, B.A.; Holman, G.M.; Wagner, R.M.; Ridgway
Biochem. Biophys. Res. Commun. 135, 622-628, 1986
A;Title: Isolation and primary structure of a peptide from the corpora cardiaca of Helic
A;Reference number: A24244; MUID:86186794; PMID:3964263
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            A;Molecule type: protein
A;Residues: 1-9 <JAF>
(S;Superfamily: adipokinetic hormone
C;Superfamily: amidated carboxyl end; corpora cardiaca; hormone; neuropeptide; pyroglutamid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
F;9/Modified site: amidated carboxyl end (Gly) #status experimental
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R;Barra, D.; Falconieri Erspamer, G.; Simmaco, M.; Bossa, F.; Melchiorri, P.; Erspamer, R;Barra, D.; Falconieri Erspamer, G.; Simmaco, M.; Bossa, F.; Melchiorri, P.; Erspamer, A;Title: Rchdei-litorin: a new peptide from the skin of Phyllomedusa rohdei.
A;Reference number: S07241; MUID:85127560; PMID:3838283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C'Superfamily: gastrin-releasing peptide
C'Keywords: amidated carboxyl end; blocked amino end; neuropeptide; pyroglutamic acid
P:1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
P:9/Modified site: amidated carboxyl end (Met) #status experimental
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C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 19-Dec-1993 #sequence_revision 25-Aug-1995 #text_change 21-Jul-2000
C;Accession: E49033; h94033
R;Morita, C.T.; Verma, S.; Aparicio, P.; Martinez, C.; Spits, H.; Brenner, M.B. Bur. J. Immunol. 21, 2999-3007, 1991
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C;Species: Phyllomedusa rohdei (Rohde's leaf frog)
C;Date: 12-Feb-1993 #sequence_revision 12-Mar-1993 #text_change 18-Aug-2000
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                                                Length 8;
                                         65.2%; Score 15; DB 2; Le
100.0%; Pred. No. 2.8e+05;
iive 0; Mismatches 0;
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Gaps

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0; Indels

0; Mismatches

Conservative

7

Matches

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C;Species: Pseudomonas aeruginosa
C;Date: 21-Jul-1995 #sequence_revision 28-Jul-1995 #text_change 21-Aug-1998
C;Accession: C53652
R;Ochsner, U.A.; Fiechter, A.; Reiser, J.
B.D.L. Chem. 269, 1997-19795, 1994
A;Title: Isolation, characterization, and expression in Escherichia coli of the Pseudomo
A;Reference number: A53652; MUID:94327521; PMID:8051059
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        phospholipase A2 (EC 3.1.1.4) precursor - sheep (fragment)
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Species: Ovis orientalis aries, Ovis ammon aries (domestic sheep)
C;Accession: A29169
R;Dutilh, C.E.; Van Doren, P.J.; Verheul, F.E.A.M.; De Haas, G.H.
Rutile: Isolation and properties of prophospholipase A2 from ox and sheep pancreas.
A;Reference number: A94661
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A;Meaidues: 1-12 <DUT>
C;Superfamily: phospholipase A2
C;Keywords: carboxylic ester hydrolase; pyroglutamic acid
F;1/Modified site: pyrrolidone carboxylic acid (Gln) #status experimental
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                                                                                                                                                                                               rhlR protein - Pseudomonas aeruginosa (fragment)
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                                                                                                                                                                                                                                                                                                                                                                                                                    A,Accession: C53652
A,Status: preliminary
A,Molecule type: DNA
Residues: 1-11 < COCH>
A,Cross-references: GB:L28170
C,Superfamily: sdiA regulatory protein
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Best Local Similarity
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February 12, 2003, 11:11:31 ; Search time 2.70833 Seconds
(without alignments)
76.572 Million cell updates/sec
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-09-403-440A-1 23 1 FXXLW 5 Title: Perfect score: Sequence:

BLOSUM62 Gapop 10.0 , Gapext 0.5

Scoring table:

112892 seqs, 41476328 residues Searched:

Total number of hits satisfying chosen parameters: Minimum DB seq length: 0 Maximum DB seq length: 25

1520

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

SwissProt_40:* Database : Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

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-	16	e,		Н	0	P82096 litoria rub	
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7	15	65.2	20	-	A1BG_EQUAS	P39090 equus asinu	
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σ.	15	65.2	22	-	CR31_LITSP		
10	15	65.2	22	Н	CR32_LITCE	litoria	
Ξ:	15	65.2	22	-	CR33_LITCE		
12	15	65.2	22	7	CR34_LITCE		
13	15	65.2	23	7	CR41_LITCE		
14	15	65.2	23	-	CR42_LITCE		
15	15	65.2	23	Н	CR43_LITCE		
16	15	65.2	23	-	SODP_PICAB		
17	15	65.2	24	7	DMS6_PHYBI		
18	15	65.2	24	7	PCL1_PACGO		
13	15	65.2	24	П	PCL2_PACGO		
20	15	65.2	24	-	VGJ_BPAL3		
21	15	65.2	24	, 1	Y3KD_NEUCR	P22702 neurospora	
55	15	65.2	25	~	VGJ_BPG4		
23	14	6.09	20	-	COXA_THUOB	P80972 thunnus obe	
24	14	6.09	24	н	COXC_THUOB	P80973 thunnus obe	
25	14	6.09	25	-	COX1_PARVE		
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SEQUENCE.
TISSUE=Ovary;
MEDLINE=94019818; PubMed=8413630;
Moore M.S., Blobel G.;
"The GTP-binding protein Ran/TC4 is required for protein import into the nucleus.";

01-0CT-1996 (Rel. 34, Created)
01-0CT-1996 (Rel. 34, Last sequence update)
01-0CT-1996 (Rel. 34, Last annotation update)
GTP-binding nuclear protein RAN (TC4) (Fragments).
RAN.
Xenopus laevis (African clawed frog).
RAN.
Manbibia; Batrachia; Anura; Mesobatrachia; Pipoidea; Pipidae;
Xenopodinae; Xenopus.

24 AA.

PRT;

STANDARD;

RAN XENLA

RESULT 2 RAN_XENLA

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P13211 P27503 P46365 P80436 P08521 P20834 P35919 P35919 P35920 P35921				Litoria rubella (Desert tree frog). Bukaryota, Metazoa, Chordata; Craniata; Vertebrata; Euteleostomi; Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae; Pelodryadinae; Litoria.	buzzing es from	(C64;	Length 6; ;; 2; Indels	
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         -!- FUNCTION: GTP-BINDING PROTEIN INVOLVED IN NUCLEOCYTOPLASMIC TRANSPORT. REQUIRED FOR THE IMPORT OF PROTEIN INTO THE NUCLEUS AND ALSO FOR RNA EXPORT. INVOLVED IN CHROMATIN CONDENSATION AND CONTROL OF CELL CYCLE (BY SIMILARITY).
-!- SUBCELLULAR LOCATION: Nuclear.
-!- SIMILARITY: BELONGS TO THE RAN FAMILY IN THE RAS SUPERFAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Rohdei-litorin: a new peptide from the skin of Phyllomedusa rohdei."; FBBS Lett. 182:53-56(1985).
-i - SUBGELLUAR: LOCATION: Secreted.
-i - SIMILARITY: BELONGS TO THE BOMBESIN/NEUROMEDIN B/RANATENSIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                     0; Gaps
                                                                                                                                                                                                                                                                                                                                                              Phyllomedusa rohdei (Rohde's leaf frog).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Phyllomedusinae; Phyllomedusa.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ·,
                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=85127560; PubMed=3838283;
Barra D., Erspamer G.F., Simmaco M., Bossa F., Melchiorri P.,
Erspamer V.;
                                                                                                                                                                      Score 16; DB 1; Length 24;
Pred. No. 9.18+02;
Trentant 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PYRROLIDONE CARBOXYLIC ACID.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       65.2%; Score 15; DB 1; Length 9; 100.0%; Pred. No. 1.1e+05;
                                                                                                                                                24 24 AA; 2657 MW; B69F83236247A250 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 4ECCC1E861ADC377 CRC64;
                                                                                                                                       GTP (BY SIMILARITY).
                                                                      HSSP, P17080, 1A2K.
InterPro, IPR002041, RAN.
PROSITE, PS01115, RAN, PARTIAL.
GTP-binding; Nuclear protein; Protein transport.
                                                                                                                                                                                                                                                                                                                     01-NOV-1988 (Rel. 09, Created)
01-FEB-1994 (Rel. 28, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 P80474;
01-OCT-1996 (Rel. 34, Created)
01-OCT-1998 (Rel. 34, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .larity 100.0%; Pred. No. 1.1. Conservative 0; Mismatches
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InterPro; IPR000074; Bombesin.
Pfam; PP02044; Bombesin; 1.
PROSITE; ES00257; BOMBESIN; 1.
Bombesin family; Amidation.
                                                                                                                                                                                69.6%;
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Best Local Similarity 40.0°
Matches 2; Conservative
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Nature 365:661-663 (1993)
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les 2; Conserv
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LITR_PHYRO
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This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the Buropean Bioinformatics Institute. There are no restrictions on its muse by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
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Biochim. Biophys. Acta 950:303-312(1988).
-!- FUNCTION: THIS PROTEIN IS INVOLVED IN CONTROL OF THE BIOSYNTHESIS
                                                                                                                                                                                                                                                                                                                                                                                                                                       MICCODIOLOGY 141:3087-3093 (1995).

-!- FUNCTION: AMINOPEPTIDASE WHICH HYDROLYZES SUBSTRATES WITH FREE N-
-!- FREMINAL AMINO ACID RESIDUES BUT NOT N-TERMINAL BLOCKED ONES.
OPTIMUM ACTIVITY IS MEASURED AT PH 7.5. MAY BE IMPORTANT IN THE
NUTRITION AND PATHOGENESIS OF THE ORGANISM IN THE HUMAN ORAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRAIN=HB8 / ATCC 27634;
MEDLINE=89000781; PubMed=2844259;
Sato S., Nakada Y., Kanaya S., Tanaka T.;
"Molecular cloning and nucleotide sequence of Thermus thermophilus
                                                                                                                                                                                                                                                                                                                                         Spratt D.A., Greenman J., Schaffer A.G.; "Capnocytophaga gingivalis aminopeptidase: a potential virulence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          .,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Thermus thermophilus.
Bacteria, Thermus/Deinococcus group, Deinococci; Thermales,
01-OCT-1996 (Rel. 34, Last annotation update)
Aminopeptidase (BC 3.4.11.-) (Fragment).
Capnocytophaga gingivalis.
Bacteria, Bacteroidetes; Flavobacteria; Flavobacteriales;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  / Match 65.2%; Score 15; DB 1; Length 10; Local Similarity 100.0%; Pred. No. 6.5e+02; nes 2; Conservative 0; Mismatches 0; Indels
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SRQUENCE 11 AA; 1228 MW; 364B295A772DC5A7 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   10 10 10 10 10 AA; 1306 MW; 00C0A6DB43772694 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -!- COFACTOR: REQUIRES MAGNESIUM OR CALCIUM.
Hydrolase; Aminopeptidase; Magnesium; Calcium.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1988 (Rel. 09, Last sequence update) 30-MAY-2000 (Rel. 39, Last annotation update) Trp operon leader peptide.
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                                                                                                                                                                                                                                    SEQUENCE.
STRAIN=AICC 33624;
MEDLINE=96118234; PubMed=8574402;
                                                                                                                                         Flavobacteriaceae; Capnocytophaga.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-NOV-1988 (Rel. 09, Created)
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NCBI_TaxID=274;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PIR; S03315; LFTWWE.
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Best Local Similarity
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                                                                                                                                                                          NCBI_TaxID=1017;
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CR31 LITSP
P56238;
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NDK_CANAL
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                                                                                                               16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last sequence update)
Peptide hormone 4 (Pea-YLS-amide).
Periplaneta americana (American cockroach).
Pukaryota, Merazoa, Arthropoda, Mandibulata, Pancrustacea, Hexapoda, Insecta, Pterygota, Neoptera, Orthopteroidea, Dictyoptera, Blattaidea, Blattaidae, Periplaneta.
                                                                                                                                                                                                                              TISSUE=Abdominal perisympathetic organs;
MEDINE=20140865; PubMed=10674456;
Predel R., Eckert M., Hollman G.M.;
"The unique neuropeptide pattern in abdominal perisympathetic organs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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  Gaps
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PREDINES=13130579; PubMed=1868686;
Patterson S.D., Bell K., Shaw D.C.;
"Donkey and horse alpha 1 B-glycoprotein: partial characterization
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Būkaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Perissodactyla, Equidae, Equus.
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-!- FUNCTION: PLASMA GLYCOPROTEIN OF UNKNOWN FUNCTION.
-!- SUBCELLULAR LOCATION: Extracellular.
-!- SIMILARITY: BELONGS TO THE IMMUNOGLOBULIN SUPERFAMILY.
Immunoglobulin domain; Glycoprotein; Plasma.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     h Similarity 100.0%; Pred. No. 1.2e+03; 2; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                       65.2%; Score 15; DB 1; Length 17; 100.0%; Pred. No. 1.1e+03; rative 0; Mismatches 0; Indels
  Indels
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17 AA; 1968 MW; CAAFB57ECD8218A9 CRC64;
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-!- FUNCTION: UNKNOWN.
-!- MASS SPECTROMETRY: MW=1966.48; METHOD=MALDI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
01-FEB-1995 (Rel. 31, Last annotation update)
Alpha-1B-glycoprotein (Postalbumin) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 AA
                                                                                            17 AA
  0; Mismatches
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                                                                                                                                                                                                                   AND MASS SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                 Neuropeptide; Amidation.
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  Conservative
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Best Local Similarity
Matches 2; Conserv
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Matches (2;
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SEQUENCE
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P39090;
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16-OCT-2001 (Rel. 40, Created)
16-OCT-2001 (Rel. 40, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Nucleoside diphosphate kinase (EC 2.7.4.6) (NDF kinase) (NDFK)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Litoria splendida (Magnificent tree frog), and
Litoria gilleni (Centralian tree frog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                  MEDIINE=96019963; PubMed=7487065;
Biondi R.M., Veron M., Walz K., Passeron S.;
"Candida albicans nucleoside-diphosphate kinase: purification and
                                                                                                                                                                                       Candida albicans (Yeast).
Eukaryota, Fungi, Ascomycota, Saccharomycotina, Saccharomycetes;
Saccharomycetales, mitosporic Saccharomycetales; Candida.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE, AND MASS SPECTROMETRY.
SPECIES-L.splendida; TISSUE-Parotoid gland;
Stone D.J.M., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. Structures of the caerins and caeridin 1 from Litoria splendida.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .;
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15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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                                                                                                                                                                                                                                                                                                                 SEQUENCE, AND CHARACTERIZATION
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STANDARD;
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Best Local Similarity
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Gaps

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Caerin 3.3.
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SEQUENCE
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SEQUENCE
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                                                                                                                                                                                                                                                                             Query Match
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                             PERCISEL. Gilleni; TISSUE-Parotoid gland;
Waugh R.J., Stone D.J.M., Bowie J.H., Wallace J.C., Tyler M.J.;
Waugh R.J., Stone D.J.M., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structures of the caerins and caeridins from Litoria gilleni.";
J. Chem. Res. 139:937-961(1993).
-!- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELLCAL CONFORMATION WHICH CAN DISRUPT BACTERIAL MEMBRANES. EACH CAERIN DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  J. Chem. Res. 138:910-936(1993).
-1- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELICAL CONFORMATION WHICH CAN DISRUPT BACTERIAL MEMBRANES. EACH CAERIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Stone D.J.M., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.; "Peptides from Australian frogs. The structures of the caerins from
                                                                                                                                                                                                                                                                                                                                                                                                                                  Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphiblai, Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria.
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                                                                                                                                                                                                               65.2%; Score 15; DB 1; Length 22; llarity 100.0%; Pred. No. 1.4e+03; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                        22 AA; 2385 MW; 1D4411E2E9D43739 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1.4e+03;

    Chem. Soc. Perkin Trans. 1:3173-3178(1992).

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15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (Rel. 36, Created)
(Rel. 36, Last sequence update)
(Rel. 40, Last annotation update)
                                                                                                                                                     -1- MASS SPECTROMETRY: MW=2382; METHOD=FAB.
Antibiotic; Amphibian skin; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        -1- MASS SPECTROMETRY: MW=2397; METHOD=FAB. Antibiotic; Amphibian skin; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ilarity 100.0%; Pred. No. 1.4
Conservative 0; Mismatches
                                                                                                                                                                               AMIDATION.
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                                                                                                                                                                                                                                                                                                                                                                                                                     Litoria caerulea (Green tree frog)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE, AND MASS SPECTROMETRY.
                       SEQUENCE, AND MASS SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                                                STANDARD;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TISSUE=Parotoid gland;
                                                                                                                                                                                                                          Local Similarity
tes 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Local Similarity es 2; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI TaxID=30344;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Litoria caerula."
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16-OCT-2001
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P56240}
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SEQUENCE
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                                                                                                                                                                                     SEQUENCE
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                                                                                                                                                                                                                                                                                                                       RESULT 10
CR32 LITCE
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CR33_LITCE
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                                                                                                                                                                                                                                                                                                   Litoria caerula.";
J. Chem. Res. 138:910-936(1993).
-!- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELICAL CONFORMATION HATCH CAN DISRUPT BACTERIAL MEMBRANES. BACH CAERIN DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Litoria caerula.";
J. Chem. Res. 138:910-936(1993).
-!- FUNCTION: ANTIBACTERIAL PREPIDE, THAT ADOPTS AN ALPHA HELICAL CONFORMATION WHICH CAN DISUPT BACTERIAL MEMBRANES. EACH CAERIN DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL
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                                                                                                                                                                                                           TISSUE=Parotoid gland;
Stone D.J.M., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structures of the caerins from
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Stone D.J.M., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.;
"Peptides from Australian frogs. The structures of the caerins from
                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Hylidae;
Pelodryadinae; Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0;
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22 AA; 2427 MW; 1D440B2200D4367C CRC64;
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22 AA; 2455 MW; 3AB40B2200D43663 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            65.2%; Score 15; DB 1; Le ilarity 100.0%; Pred. No. 1.4e+03; Conservative 0; Mismatches 0;
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16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -1- MASS SPECTROMETRY: MW=2424; METHOD=FAB.
Antibiotic; Amphibian skin; Amidation.
MOD RES 22 AMIDATION.
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Litoria caerulea (Green tree frog).
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                                                                                                                                                                                       SEQUENCE, AND MASS SPECTROMETRY.
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es 2; Conserv
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Best Local Similarity
                                                                                                                            NCBI_TaxID=30344;
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CR43 LITCE
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J. Chem. Res. 138:910-936(1993).
-!- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELICAL CONFORMATION WHICH CAN DISRUPT BACTERIAL MEMBRANES. EACH CAERIN DISPLAYS A DIFPERRNY ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL.
                                                                                                                                                                                                                                                                                                                                                                                                                         Litoria caerula.";
J. Chem. Res. 138:910-936(1993).
-!- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELICAL CONFORMATION HAICH CAN DISRUPT BACTERIAL MEMBRANES. EACH CAERIN DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
-!- TISSUE SPECIFICITY: SECRETED BY THE SKIN PAROTOID AND/OR ROSTRAL.
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Amphibia; Batrachia, Anura; Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria.
                                                                                                                                                        Litoria caerulea (Green tree frog).
Skaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                65.2%; Score 15; DB 1; Length 23; 100.0%; Pred. No. 1.4e+03; tive 0; Mismatches 0; Indels
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15-UUL-1998 (Rel. 36, Created)
15-UUL-1998 (Rel. 36, Last sequence update)
116-CCT-2001 (Rel. 40, Last annotation update)
Caexin 4.2.
                                                 15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
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  23 AA.
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  PRT;
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                                                                                                                                                                                                                                                                                                                   SEQUENCE, AND MASS SPECTROMETRY. TISSUE=Parotoid gland;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE, AND MASS SPECTROMETRY.
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Matches 2; Conservative
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  STANDARD;
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CR41 LITCE
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  HAD DESCRIBED TO SERVING SECOND SECON
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4 LW 5

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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                     Litoria caerula.";
J. Chem. Res. 138:910-936(1993)
J. Chem. Res. 138:910-936(1993)
-!- FUNCTION: ANTIBACTERIAL PEPTIDE, THAT ADOPTS AN ALPHA HELICAL
CONFORMATION WHICH CAN DISRUPT BACTERIAL MEMBRANES. BACH CAERIN
DISPLAYS A DIFFERENT ANTIMICROBIAL SPECIFICITY.
                                                                                                                                                                                                                                                                                                                                                    Stone D.J.M., Waugh R.J., Bowie J.H., Wallace J.C., Tyler M.J.; "Peptides from Australian frogs. The structures of the caerins from
                                                                                                                                                                          Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Hylidae,
Pelodryadinae, Litoria.
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Pred. No. 1.4e+03;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23 AA; 235 AMIDATION.
23 AA; 2356 MW; B98FDD80B6ADDC9D CRC64;
                                                 15-JUL-1998 (Rel. 36, Created)
15-JUL-1998 (Rel. 36, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           65.2%; Scor.
100.0%; Pred. No. 1...
     23 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Search completed: February 12, 2003, 11:14:05 Job time : 3.70833 secs
                                                                                                                                                       Litoria caerulea (Green tree frog)
                                                                                                                                                                                                                                                                                                       SEQUENCE, AND MASS SPECTROMETRY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
     STANDARD;
                                                                                                                                                                                                                                                                                                                                 TISSUE=Parotoid gland;
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les 2; Conserv
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CR43 LITCE
P56244;
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SEQUENCE
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Pebruary 12, 2003, 11:12:02 ; Search time 9.79167 Seconds
(without alignments)
105.216 Million cell updates/sec
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GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                                  Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                 671580 segs, 206047115 residues
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Maximum Match 100%
Listing first 45 summaries
                                                                     OM protein - protein search, using sw model
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1: Sp archea:*
2: Sp_bacteria:*
3: Sp_fungi:*
4: Sp_fungi:*
5: Sp_fungi:*
5: Sp_man:*
5: Sp_mhc:*
5: Sp_mhc:*
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Gapop 10.0 , Gapext 0.5
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23
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Perfect score:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. sp_archeap:* 11123

sp_rodent:*
sp_virus:*
sp_vertebrate:*
sp_unclassified:*

sp_bacteriap:*

rvirus:*

	Description	09w6d7 gallus gall	Ogno68 homo sapien	O8uvw3 brachydanio	O61943 mus musculu	O9bvm6 homo sapien	053579 rhodobacter	053580 rhodobacter	O9dw76 mus sp. hom	035353 rattus norv	Q66548 human herpe	Q72017 human immun	O9s0u6 shiqella so	P72429 salmonella	085406 coxiella bu	095953 homo sapien	Q38366 bacteriopha
SUMMARIES	ΩI	09W6D7	O9NP68	1 QBUVW3	961943	Q9BYM6	Q53579	Q53580	94WQ6Q	. 035353	. Q66548	. Q72017	900860	P72429	085406	095953	038366
	igth DB	22 13	25 4	25 13	21 11	23 4	12 2	15 2	16 11	20 11	20 12	17 15	25 2	25 16	8	9	6
ф	Query Match Length DB	87.0	87.0	87.0	82.6	78.3	73.9	73.9	73.9	73.9	73.9	9.69	9.69	9.69	65.2	65.2	65.2
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25 AA.

O9NP68 PRELIMINARY; PRT; O9NP68; 01-OCT-2000 (TrEMBLrel. 15, Created)

RESULT 2 Q9NP68 ID Q9 AC Q9 DT 01

Q8wtt4 homo sapien Q8shf6 chamaeleo m Q9esu5 mus musculu Q47602 escherichia Q9ucr1 homo sapien Q9ucr1 homo sapien Q9g148 sus scrofa O77895 oreochromis Q9g148 cus scrofa O77895 oreochromis Q9g204 otcorryptis Q9379 rattus norv Q98240 human immun Q42230 mycoplasma Q3x12 bemisia tab Q3	Tudels
65.2 10 4 08WTT4 65.2 10 8 08SHF6 65.2 10 10 10 09ESU5 65.2 11 4 09UG81 65.2 11 4 09UG81 65.2 11 7 077895 65.2 11 7 077895 65.2 11 7 077895 65.2 12 10 063579 65.2 12 10 063579 65.2 12 10 063579 65.2 12 10 063579 65.2 12 10 063579 65.2 14 6 077538 65.2 14 8 09WTG1 65.2 14 10 09SUF8 65.2 14 10 09SUF8 65.2 14 10 09SUF8 65.2 14 10 09SUF8 65.2 16 13 08UUF8 65.2 14 15 09UUF8 65.2 16 13 08UUF8 65.3 16 13 08UUF8 65.3 16 13 08UUF8 65.3 16 13 08UUF8 65.3 14 08UUF8 65.3 16 13 08UUF8 65.3 14 08UUF8 65.3	; conservative v; mismatches z; W 5

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RESULT 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Actinopterygii, Neopterygii, Teleostei, Ostariophysi, Cypriniformes,
Cyprinidae, Danio.
NCBI_TaxID=7955,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Gaps
                                                                        Eukaryoča, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                               A Thompson-Hehir J., Davies M.P.A., Green J.A., Halliwell N.,
Thompson-Hehir J., Davies M.P.A., Green J.A., Halliwell N.,
A Joyce K.A., Salisbury J., Sibson D.R., Vergote I., Walker C.;
Submitted (NOV-1999) to the EWEL/GenBank/DDBJ databases.
R EMBL; AF209136; AAF36362.1; -.
R EMBL; AF209131; AAF36354.1; -.
R EMBL; AF209131; AAF36356.1; -.
R EMBL; AF209131; AAF36356.1; -.
R EMBL; AF209131; AAF36359.1; -.
R EMBL; AF209134; AAF36350.1; -.
R EMBL; AF209134; AAF36360.1; -.
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                                                                                                                                                                                                                                                                                                                                                                                                           87.0%; Score 20; DB 4; Length 25; 60.0%; Pred. No. 6.7e+02; tive 0; Mismatches 2; Indels
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NON TER 25 25
SEQUENCE 25 AA; 2837 MW; 9EB8AA36B50DB17E CRC64;
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01-WAR-2002 (TrEMBLrel. 20, Last sequence update)
01-UN-2002 (TrEMBLrel. 21, Last annotation update)
AMPA receptor subunit 2 (Fragment).
01-OCT-2000 (TrEMBLrel. 15, Last sequence update) 01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Brachydanio rerio (Zebrafish) (Zebra danio).
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InterPro; IPR001320; Ion glu_receptor.
ProDom; PD000500; Ion_glu_receptor; 1.
                            rumor suppressor p53 (Fragment).
                                                                                                                                                                                                                                                                                                                                   InterPro; IPR002117; P53.
Pfam; PF00870; P53; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                           Best_Local Similarity 60.0
Matches 3; Conservative
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                                                           Homo sapiens (Human)
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                                                                                                          NCBI_TaxID=9606;
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Caput D.; "The gene coding for the interferon-inducible human dsRNA adenosine deaminase is transcribed into several messengers specifying different
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Neuroscience 69:807-819(1995).
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                                                                                                                                                                                                                                                            Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Rodentia, Sciurognathi, Muridae, Murinae, Mus.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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0
                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE=96164254; PubMed=8596650;
Bessis A., Salmon A.M., Zoli M., Le Novere N., Picciotto M.,
Changeux J.P.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  h 82.6%; Score 19; DB 11; Length 21; Similarity 60.0%; Pred. No. 9.3e+02; 3; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bessis A.E.;
Submitted (JUL-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; X82655; CAA57978.1; -.
MGD; MGI:87891; Chrnb2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        proteins.";
Submitted (JUN-1996) to the EMBL/GenBank/DDBJ databases.
EMBL; X98560; CAA67171.1; -.
NON TER 23 .... ww. CP09179B7DFC1395 CRC64;
                                                    01-NoV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 19, Last annotation update)
01-DBC-2001 (TrEMBLrel. 19, Last annotation update)
Nicotinic acetylcholine receptor beta 2-subunit protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Receptor.
NON TER 21 21
SEQÜENCE 21 AA; 2351 MW; 28E57E2BC2BC67FA CRC64;
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01-JUN-2001 (TrEMBLrel. 17, Last sequence update)
01-JUN-2001 (TrEMBLrel. 17, Last annotation update)
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PRELIMINARY;
                                                                                                                                                                                                                                      Mus musculus (Mouse)
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                                                                                                                                                                                 (Fragment).
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Q61943
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Bacteria; Proteobacteria, alpha subdivision; Rhodobacter group;
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Bacteria; Proteobacteria; alpha subdivision; Rhodobacter group;
Rhodobacter.
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MEDILTE-22234963; bubMed=1569029;
Richter P., Brand M., Drews G.;
"Characterization of LHI- and LHI+ Rhodobacter capsulatus pufA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
MEDLINE-9224963; bubmed=1569029;
Richter P., Brand M., Drews G.;
"Characterization of LHI- and LHI+ Rhodobacter capsulatus pufA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              h Similarity 40.0%; Pred. No. 1.8e+03; Length 15; 2; Conservative 1; Mismatches 2; Indels
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                                                                                                                                                                                                                             01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1996 (TrEMBLrel. 19, Last annotation update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Light-harvesting complex I alpha polypeptide (Fragment)
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
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EMBL; S97552; AAC60406.1; -.
NON TER 15
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J. Bacteriol. 174:3030-3041(1992).
EMBL; S97551; AAC60405.1; -.
NON TER 12. 12. 12.
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                                               19 FAAVW 23
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1 FXXLW 5
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Q53580;
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Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A., AND DISTRIBUTION IN BRAIN.
STRAIN=SPRAGUE-DAWLEY: TISSUE=BRAIN;
MEDLINE=98281588; PubMed=9622245;
MEDLINE=98281588; M.A. Frandes K.J., Cockett M.I.;
"Distribution of heterotrimeric G-protein beta and gamma subunits in
                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Guanine nucleotide-binding protein beta subunit 4 (Transducin beta
                                                                                                   Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Mus.
NCBI_TaxID=10095;
                                                                                                                                                                                                                                                                                                                                                                                                                ;
                                                                                                                                                                      SEQUENCE FROM N.A.

SEQUENCE FROM N.A.

SIGNIBLINE-22073356; PubMed=1683707;

Singh G., Kaur S., Stock J.L., Jenkins N.A., Gilbert D.J.,

Copeland N.G., Potter S.S.;

Tidentification of 10 murine homeobox genes.";

Proc. Natl. Acad. Sci. U.S.A. 88:10706-10710(1991).

InterPro; IPR001356; Homeobox.
                                                                                                                                                                                                                                                                                                                                                                             73.9%; Score 17; DB 11; Length 16; 40.0%; Pred. No. 1.9e+03; ative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                  16 AA; 2183 MW; BE902130A4525041 CRC64;
                          01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Homeobox GSH-3 (Fragment).
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NON TER <1 20 WD7.
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PROSITE; PS50071; HOMEOBOX_2; 1.
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PRELIMINARY;
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RESULT 10

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Conservative
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                      PRELIMINARY;
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                                                                                                      ORF10S (Fragment).
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tes 2; Conserv
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                                                                                                                     Shigella sonnei
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MEDLNE=88217505; PubMed=2835748;
Walls D., Perricaudet M., Gannon F.;
"The analysis of BBV proteins which are antigenic in vivo.";
Nucleic Acids Res. 16:2859-2872(1988).
EMBL; X07531; CAA30406.1;
NON TER
                             / Match 73.9%; Score 17; DB 11; Length 20; Local Similarity 40.0%; Pred. No. 2.4e+03; nes 2; Conservative 1; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 73.9%; Score 17; DB 12; Length 20; Best Local Similarity 40.0%; Pred. No. 2.4e+03; Matches 2; Conservative 1; Mismatches 2; Indels
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                                                                                                                                                                                                                                           01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
EBV B95-8 Cl(g) DNA with antigen-coding ORF (Fragment) whuman herpesvirus 4 (Eppetain-Barr virus).
Viruses, daDNA viruses, no RNA stage; Herpesviridae;
dammaherpesvirinae; Lymphocryptovirus.
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Michael N.L.;
Submitted (APR-1995) to the EMBL/GenBank/DDBJ databases.
EMBL; U24486; AAA79676.1; -.
NON_TER.
20 AA; 2242 MW; C143688295AE0BE0 CRC64;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Human immunodeficiency virus type 1.
Viruses; Retroid viruses; Retroviridae; Lentivirus.
NCBI_TaxID=11676;
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Best Local Similarity 40.03
Matches 2; Conservative
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11 FRFIW 15
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RESULT 11

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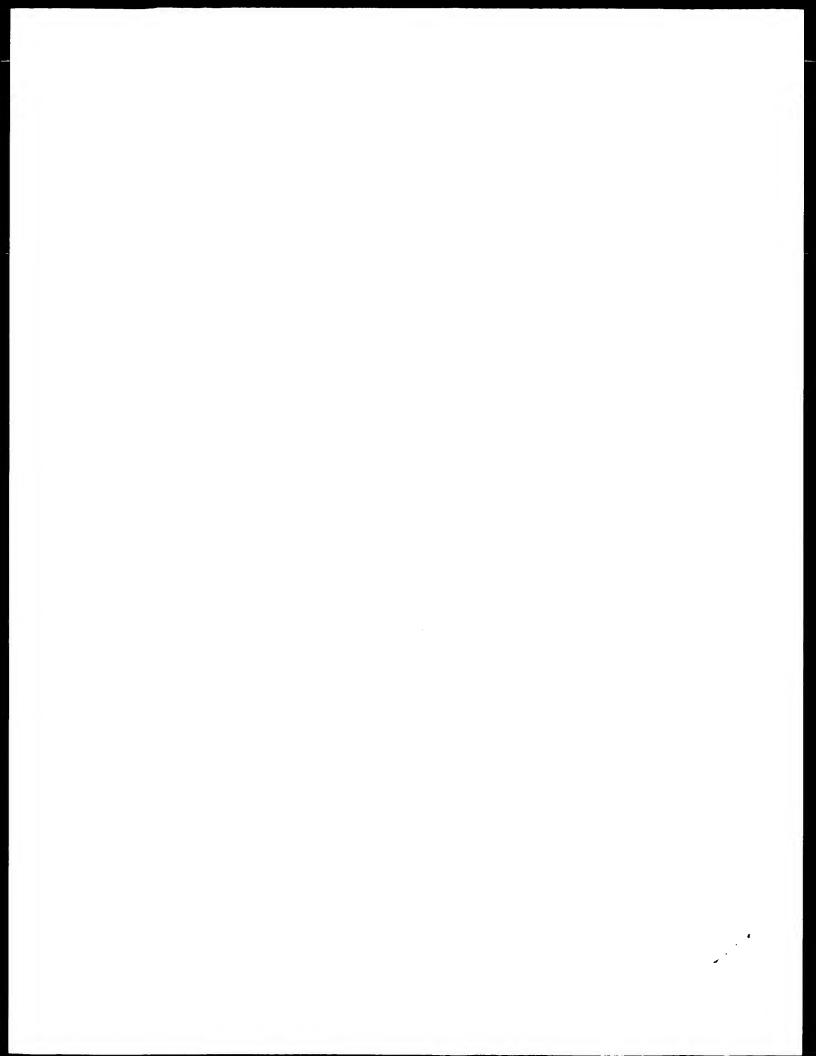
RESULT 12

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STRAIN=HW383; MEDLINE=20248365; PubMed=10789503; MEDLINE=20248365; PubMed=10789503; Chida T., Okamra N., Ohtani K., Yoshida Y., Arakawa E., Watanabe H.; "The complete DNA sequence of the O antigen gene region of Plessomon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Houng H.H., Venkatesan M.M., "Genetic analysis of Shigella sonnei form I antigen: identification of a novel 18630 as an essential element for the form I antigen
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                                                                                                                       Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
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"50 kb virulence plasmid of Salmonella enterica serovar
Choleraesuis.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. STRAIN=82139; PLASMID=VIRULENCE; SPECIES=S.enteritidis; STRAIN=82139; PLASMID=VIRULENCE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A Microb. Pathog. 25:165-173(1998).
Microb. Pathog. 25:165-173(1998).
EMBL; AB028134; BAA85070.1; -.
NON TER 25 25
SEQUENCE 25 AA; 2726 MW; A6CC5D24211CDAAD CRC64;
01-MAY-2000 (TrEMBLrel. 13, Created)
01-MAY-2000 (TrEMBLrel. 13, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-FEB-1997 (TrEMBLrel. 02, Created)
01-FEB-1997 (TrEMBLrel. 02, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
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Microbiol. Immunol. 44:161-172(2000).
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                                                                                SPECIESES. Typhimurium; STRAIN=LT2 / SGSC1412 / ATCC 700720; PLASMID=PSLT; PLASMID=PSLT; PLASMID=PSLT; PLASMID=PSLT; PLASMID=PSLT; PLASMID=PSLT; MCDELLand M., Sanderson K.E., Spieth J., Clifton S.W., Latreille P., Courtney L., Porwollik S., Ali J., Dante M., Du F., Hou S., Layman D., Leonard S., Mauyen C., Scott K., Holmes A., Grewal N., Mulvaney B., Ryan B., Sun H., Florea L., Miller W., Stoneking T., Nhan M., Waterston R., Wilson R.K.; "Complete genome sequence of Salmonella enterica serovar Typhimurium
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-NOV-1998 (TrEMBLrel. 19, Last sequence update)
01-DEC-2001 (TrEMBLrel. 19, Last annotation update)
Hypothetical 1.0 kDa protein (Fragment).
Coxiella burnetii.
Bacteria; Proteobacteria; gamma subdivision; Legionellaceae group;
Coxiella group; Coxiella.
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MEDLINE=98348442; PubMed=9683477;
Willems H., Jaeger C., Baljer G.;
Willems H., Jaeger C., Baljer G.;
Willems H., Jaeger C., Baljer G.;
J. Bacteriol and genetic map of the obligate intracellular bacterium coxiella burnetii.";
J. Bacteriol 180:3816-3822(1998).
Hypothetical protein.
NON_TER
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Pred. No. 4.8e+03;
1; Mismatches 2; Indels
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Submiveed (MAR-2000) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; U64797; AAB53038.1; -.
EMBL; AB040415; BAB20531.1; -.
EMBL; AE006471; AAL23447.1; -.
Plasmid; Complete proteone.
SEQUENCE 25 AA; 3008 WW; B21F596B6DEFABE7 CRC64;
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-MAY-2002 (TrEMBLrel. 20, Last annotation update)
Galactocerebrosidase (EC 3.2.1.46) (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                   Nature 413:852-856(2001)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
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                                                            SEQUENCE FROM N.A.
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085406
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095953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Matches
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REPRESENT SERVICE SERV
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OC Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; OC Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
OX NCBL TaxID=9606;
RN [1]
RP SEQUENCE FROM N.A.
RA Lulli L., Torchiana E., Finocchiaro G.;
RR Lulli L., Torchiana E., Finocchiaro G.;
RR Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
RE SUBMIT AAD15626.1; -..
KW Glycosidaes; Hydrolase.
FT NON TER 9
SQUENCE 9 AA; 1069 MW; 374E2AADC2C699C8 CRC64;
Ouery Match
Best Local Similarity 100.0%; Pred. No. 6.7e+05;
Matches 2; Conservative 0; Mismatches 0; Indels 0; Gaps
OY 4 LW 5
Db 8 LW 9
Search completed; February 12, 2003, 11:14:58
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February 12, 2003, 11:10:52; Search time 12.5 Seconds (without alignments) 53.300 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                               320064
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                         908470 seqs, 133250620 residues
                                                                                                                                                                                                                                                                                                                                                                                                                              Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                      OM protein - protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                              US-09-403-440A-1
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Perfect score:
                                                                                                                                                                                                                                                    Scoring table:
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                                                                                                                                                                                                                  Sequence:
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution. /SIDS2/gcgdata/geneseq/geneseqp-embl/AA1999.DAT:*/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2000.DAT:*/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2001.DAT:*/SIDS2/gcgdata/geneseq/geneseqp-embl/AA2002.DAT:*

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. DAT: *

| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1980.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1981.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1987.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1987.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1987.DAT:*
| SIDS2/gcgdata/geneseq/geneseqp-embl/AA1987.DAT:*

			Description	Peptide #8. Unide	Carbohydrate antiq	Motif binding to a	p53/MDM2 binding i	p53 minimal MDM2 b	p53 protein amino	Mdm/hdm antagonist	Mdm/hdm antagonist	Antibody DO-1 epit	p53 binding site k
SUMMARIES			E	AAB29164	AAY33038	AAB07280	AAR89913	AAR89915	AAW13606	AAB17074	ABB73169	AAR89921	AAR89925
			DB	21	20	21	17	17	18	21	23	17	17
	de		Match Length DB	15	S	ß	9	9	9	9	9	7	7
		Query	Match	91.3	87.0	87.0	87.0	87.0	87.0	87.0	87.0	87.0	87.0
			Score	21	20	20	20	20	20	20	20	20	20
		Result	No.	7	7	m	4	ß	9	7	60	თ	10

Assays and screening methods based on direct interaction between FHA domains and phosphopeptides, useful for characterizing binding and to identify binding partners and modulators of FHA domain-phosphopeptide

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ALIGNMENTS

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Fork head associated, FHA; domain, transcriptional control, DNA replication, DNA repair; cell cycle control.
              AAB29164 standard; Peptide; 15 AA
                                                                                                                                               17-MAR-2000; 2000WO-GB01024.
                                                                                                                                                               99GB-0006432.
                                              (first entry)
                                                                                                                                                                                                     Jackson SP, Durocher D;
                                                                                                                                                                                     (KUDO-) KUDOS PHARM LTD.
                                                                                                                                                                                                                    WPI; 2000-664872/64.
                                                                                                                 WO200057184-A2
                                                                                                                                                              19-MAR-1999;
28-JUN-1999;
                                                                                                  Unidentified
                                                                                                                               28-SEP-2000.
                                             02-FEB-2001
                                                             Peptide #8.
                              AAB29164;
RESULT 1
        AAB29164
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This invention describes a novel method for preparing a peptide or recombinant antibody, which mimics an antigenic carbohydrate. The peptides and recombinant antibodies prepared to mimic antigenic carbohydrates can be used to enhance binding of anti-antigenic carbohydrates can be used to enhance binding of anti-antigenic carbohydrate antibodies to the antigenic carbohydrate, e.g. as vaccine adjuvants. The peptides can be used to inhibit binding of a ligand to a receptor, which is an antigenic carbohydrate. The methods are used to prepare the peptides and antigenic antibodies, which mimic the antigenic carbohydrates. Carbohydrate-conjugate vaccines could be used in tumour immunity and cancer therapy. The peptides and antibodies can also be used as antibacterials. Peptides that mimic carbohydrate antigens can be formulated to develop a longer lasting immune response. Other advantages
                                                                                                                                     molecules. FHA peptides may be useful for treating medical conditions associated with defects in transcriptional control, DNA replication, DNA replication, and repair, cell cycle control or other cellular process. The method may provide valuable insights into checkpoint signalling, has important implications for the functions of other FHA domain-containing proteins and provides basis for new lines of therapy. The present sequence is a peptide used in the present invention.
                                                                          The present invention relates to assays and screening methods based on a direct interaction between fork head associated (FHA) domains and phosphorylated polypeptides, for characterizing the binding of these
                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mimotope; antigen; carbohydrate; antibody; mimic; vaccine; adjuvant; tumour immunity; cancer therapy; antibacterial; immune response; immunogenicity; anti-idiotype; T cell response manipulation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Peptides and recombinant antibody mimics of carbohydrate antigens, used for, e.g. treatment of cancer and infection
                                                                                                                                                                                                                                                                                                                                                  91.3%; Score 21; DB 21; Length 15; 60.0%; Pred. No. 5.3e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Carbohydrate antigen peptide mimotope 11.
                                     Disclosure, Fig 2, 92pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAY33038 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 21; Page 72; 88pp; English.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-NOV-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 60.0%
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (UYPE-) UNIV PENNSYLVANIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1999-527317/44.
                                                                                                                                                                                                                                                                                                            15 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Kieber-Emmons T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             9 FSALW 13
                                                                                                                                                                                                                                                                                                                                                                                                                                   1 FXXLW 5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    04-FEB-1998;
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                                                                                                                                                                                                                                                                                                               Sequence
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tumour, merasass to take the account of marians and attachment to the endothelium of distant organs, ultimately resulting in the formation of new tumours. The selectin family of adhesion molecules is implicated in the process. E-selectin is a calcium-dependent molecule expressed by activated vascular endothelium. E-selectins bind to glycoconjugates carrying a termanal tetrasaccharide Lewis antigen, which are found on tumour cell surfaces. One such Lewis antigen, which are found on important step in metastasis. Therefore, inhibition of E-selectin-dependent carbohydrate-mediated interactions is thought to be a target for anti-cancer therapy. The present sequence is a non planar-X-planar type motif. This motif is thought to bind to anti-Ley cantibody BRS5-2. BRS5-2 is a peptido-mimetics of E-selectins. Peptides that block E-selectin-Ley binding and therefore adhesion of tumour cells and leukocytes to endothelial cells inhibit metastasis.
                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Tumour metastasis requires detachment of malignant cells from the primary
                                                                          amount of antigen required for immunization; (3) peptide synthesis may be more practical than synthesis of carbohydrate-protein conjugates or the production of anti-idiotypes; (4) peptide mimicking sequences can be engineered into DNA plasmids for DNA vaccination to further manipulate T cell responses. AAY33028-Y33087 represent the carbohydrate antigen peptide mimotopes described in the invention.
of the peptide mimics are; (1) the chemical composition and purity of synthesized peptides can be precisely defined; (2) the immunogenicity of the peptides can be significantly enhanced by polymerization or addition or relatively small carrier molecules that reduce the total
                                                                                                                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Peptidomimetics of carbohydrate Lewis ligands useful for modulating inflammation, metastasis and angiogenesis
                                                                                                                                                                                                                                                                                                                      .,
                                                                                                                                                                                                                                                                            Query Match 87.0%; Score 20; DB 20; Length 5; Best Local Similarity 60.0%; Pred. No. 7.8e+05; Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Human; peptido-mimetic; tumour metastasis; E-selectin;
adhesion molecule; Lewis antigen; anti-adhesion therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Motif binding to anti-Lewis antigen anibody BR55-2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (WIST-) WISTAR INST ANATOMY & BIOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Blaszczyk-Thurin M, Kieber-Emmons T;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Example 5; Page 37; 107pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAB07280 standard; peptide; 5 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   99WO-US26277.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          WPI; 2000-376309/32.
                                                                                                                                                                                                                                        5 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            06-NOV-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                            1 FSLLW 5
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                                                                                                                                                                                                                                        Sequence
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19-APR-1995;
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                                                                                                                                                                                   20-JUL-1994;
                    10-SEP-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              16-JAN-1998
                                                                                                                                  01-FEB-1996.
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                                                                                         Synthetic.
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AAR89915;
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                                                                                                                                                                                                                         Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New peptides of the invention which interfere with the binding of the p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptides equences ARR89911-3 which form part of the p53 protein binding motif between amino acids 16-33 (AAR89914). The peptides were identified by modifying the p53 consensus binding sequence by substitution of an Alamine at each pos. and identifying which amino acid changes altered binding to expressed MDM2. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                 New cpds. which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and
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0
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0
                         87.0%; Score 20; DB 21; Length 5; 60.0%; Pred. No. 7.8e+05; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87.0%; Score 20; DB 17; Length 6; 60.0%; Pred. No. 7.8e+05; ive 0; Mismatches 2; Indels
                                                                                                                                          AAR89913 standard; peptide; 6 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 4; Page 29; 46pp; English.
                                                                                                                                                                                                     p53/MDM2 binding inhibitor #3
                                                                                                                                                                                                                                                                                                                     95WO-GB01719
                                                                                                                                                                                                                                                                                                                                        95US-0424957.
                                                                                                                                                                                 (first entry)
                                                3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                          Lane DP, Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-105905/11.
                                                                                                                                                                                                                                                                                                                                                                     (UYDU-) UNIV DUNDEE.
                                       Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                      other malignancies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
         s AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   6 AA;
                                                                    1 FXXLW 5
                                                                                       1 FSLLW 5
                                                                                                                                                                                 10-SEP-1996
                                                                                                                                                                                                                                                                                                                                        19-APR-1995;
20-JUL-1994;
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AAR89915
ID AAR8
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                                                                                                                     RESULT 4
                                                                                                                                AAR89913
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AAR89915 standard; peptide; 6 AA.

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This is the sequence of the p53 tumour suppressor protein minimal MDM2 oncogene protein binding site. The sequence is used to generate peptides (ARR89911-3) conteg. this sequence, which forms part of the p53 protein binding motif between amino acids 16-33 (ARR89914). The peptides were identified by modifying the p53 consensus binding sequence by substitution of an Alanine at each pos, and identifying which amino acid changes altered binding to expressed MDM2. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Mouse; Mdm2; murine double minute; phosphoprotein; binding; modulation; tumour suppressor; p53; oncogene; cell cycle arrest; p107; antagonist; inhibition; transcription factor; adenocarcinoma; colon; cancer; breast; lung; stomach; myeloid leukaemia; lymphoma; hyperproliferative;
                                                                                                       Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New cpds, which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and other malignancies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 15; 46pp; English.
                                                   p53 minimal MDM2 binding site #1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAW13606 standard; peptide; 6 AA.
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                                                                                                                                                                                                                                                                                                                                                              95WO-GB01719.
                                                                                                                                                                                                                                                                                                                                                                                                               95US-0424957.
(first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity 60.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Lane DP, Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WPI; 1996-105905/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (UYDU-) UNIV DUNDEE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO9709343-A2
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The peptides AAW13602-6 represent peptide fragments derived from the wild type human p53 protein. This peptide corresponds to amino acids 18-23 of the p53 sequence. The peptides are claimed peptides which are able to bind the N-terminal amino acids (1-134) of the mutine double minute-2 (mdm2) protein (AAW13600). Mdm2 protein is a 90 kD composed protein which binds and modulates the activity of the tumour suppressor protein p53. It has now been shown that the mdm2 protein cuppressor protein properties, especially in a p53-null background. Mdm2 is observed to unblock cell cycle arrest in G1 caused by over-expression of the p107 protein. The p53 peptides are examples of antagonists of the invention which are able to inhibit the oncogenic activity of mdm2. The antagonists are used to treat e.g. adenocarcinoms of the colon, cancer of the breast, lung or stomach, myeloid leukaemia; B cell lymphoma, or other hyperproliferative conditions such as restenosis. Note: this sequence is not given in the specification but is constructed from the wild type human p53 sequence.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                autoimmune disease; crostatic; antiaschmatic; thrombolytic; VEGF; immunosuppressive; EPO; TPO; CTLA4; mimetic; IL-1; TNF; antagonist; MMP; inhibitor; erythropoietin; thrombopoietin; interleukin 1; cytotoxic T cell lymphocyte antigen 4; tumour necrosis factor; vascular endothelial growth factor; matrix metalloproteinaes; asthma; thrombosis; pharmaceutical.
                                                                                                                                                                                                                                                 Treating cancer with antagonist of oncogenic activity of protein Mdm2 - or nucleic acid encoding an antagonist, also viral vectors
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Modified peptide; therapeutic agent; fusion; Fc domain; cancer;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  87.0%; Score 20; DB 18; Length 6; 60.0%; Pred. No. 7.8e+05; 1ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mdm/hdm antagonist peptide sequence SEQ ID NO:130.
                                                                                                                                                                         Dubs-Poterszman M, Tocque B, Wasylyk B;
                                                                                                                (INRM ) INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB17074 standard; Peptide; 6 AA.
                                                                                                                                  (RHON ) RHONE POULENC RORER SA.
                                                                                                                                                                                                                                                                                                                             Claim 4; Page -; 43pp; French.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            99WO-US25044.
                                      96WO-FR01340.
                                                                           95FR-0010331.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 60.0%
Marches 3, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          31-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                      contg. this nucleic acid
                                                                                                                                                                                                            WPI; 1997-192837/17.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WO200024782-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 FXXLW 5
                                      02-SEP-1996;
                                                                           04-SEP-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 FSDLW 6
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  13-MAR-1997.
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The present invention describes composition of matter (1) comprising an Fc domain, pharmacologically active peptides, and linkers. Where (I) is: (X1)3-F1-(X2)b, where: F1 = an Fc domain; X1 and X2 = are each independently selected from -(IJ)-C-F1. (L2)d-F2.

-(L1)c-P1. (L2)d-P2-(L3)e-P^3, or -(L1)c-P1. (L2)d-P2-(L3)e-P3. (L4)f-P4 where P1, P2, P3, and P4 = are each independently sequences of pharmacologically active peptides; L1, L2, L3, and L4 = are each independently inkers; and a, b, c, d, e, and f = are each independently inkers; and a, b, c, d, e, and f = are each independently of or 1, provided that at least 1 of a and b is 1. The composition can have cytostatic, antiasthmatic, thrombolytic and immunosuppressive activities. DNAs, vectors and host cells from the present invention can bave cytostatic antiasthmatic, thrombolytic and immunosuppressive activities. DNAs, vectors and host cells from the present invention can be used for producing pharmaceutical compositions are useful for treating cancer, asthma, thrombosis, or autoimmune diseases. The use of an Fc domain (rather than a Fab domain) can provide a longer half-life or incorporate functions such as Fc receptor binding, prordering complement fixation, and possibly placental transfer. AAA69443 to AAA6956 and AAB16955 to AAB18003 represent nucleotide and amino acid sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EPO; erythropoletin; TPO; tumour necrosis factor alpha inhibitor; TNF-alpha inhibitor; Interleukin lantagonist; IL-1 antagonist; TNF-alpha inhibitor; interleukin lantagonist; IL-1 antagonist; TNF); TPO mimetic peptide; EPO mimetic peptide; EPO mimetic peptide; EPO mimetic peptide; Interleukin lantagonist; MMP inhibitor; antiinflammatory; antitumour; immunosupressive; cytostatic; antiinflamthitic; antidabetic; ophthalmological; antidamemic; antimeriality; haemostatic; dermatological; neuroprotective; inflammatory disease; autoimmune disease; tumour growth; cancer; rheumatord arthritis; diabetic retinopathy; infertility; obesity; sleep disorder; neurological degenerative disease; hamemia; thrombocytopaenia; metastatic tumour; systemic lupus erythematosus;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Modified peptide; mimetic; Fc domain; fusion; immunoglobulin G; IgG;
                                                                                                                                                                                                 Novel composition of matter comprising an Fc domain and pharmacologically active peptides, useful for treating cancer and autoimmune diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               87.0%; Score 20; DB 21; Length 6; 60.0%; Pred. No. 7.8e+05; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mdm/hdm antagonist peptide SEQ ID NO:130.
                                                                                                                  Liu C, Cheetham J, Boone TC;
                                                                                                                                                                                                                                                                                                               Claim 39; Page 240; 608pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABB73169 standard; Peptide; 6 AA.
98US-0105371.
99US-0428082.
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                                                                                                                                                                WPI; 2000-350702/30.
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                                                                    (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         6 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WO200183525-A2.
23-OCT-1998;
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                      22-OCT-1999;
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                                                                                                                     Peige U,
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AAR8992!
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                                                                                                                                                                                                                                                                                                                                                       multimers. (I) can have antiinflammatory, antitumour, immunosuppressive, cytostatic, antirheumatic, antiarthritic, antitumour, immunosuppressive, cytostatic, antirheumatic, antiarthritic, antidiabetic, ophthalmological, antianamenic, anorectic, antiinfertility, heamostatic, dermacological, corporporective activities. (I) can be used as a therapeutic or prophylactic agent as well as for screening purposes. (I) is useful for diagnosing diseases characterised by dysfunction of their associated protein of interest, for identifying normal or abnormal proteins of interest in a biological sample. Additionally, (I) is useful for treating inflammatory and autoimmune diseases, tumour growth, cancer, retumatoid arthritis, diagnetic retinopathy, obesity, sleep disorders, infertility, and neurological degenerative diseases. (I), comprising low red blood cell levels such as anaemia. The TPO-mimetic comprising computed in the time of the involve an existing computed the contractive of the contractive 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 compounds are asian. The friction of an expected megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet deficiency or an expected megakaryocyte/platelet deficiency, such as thrombocytopaenia, aplastic anaemia, metastatic tumour which result in thrombocytopaenia, systemic lupus erythematosus, and Fanconi's syndromen. ABB72403 to ABB73405 and ABL35695 to ABL35777 represent amino acid and nucleic acid sequences used in the exemplification of the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus; antibody.
                                                                                                                                                                                                                                                                                                                                               The present invention describes a vehicle-peptide molecule (I) or its
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                              Novel vehicle-peptide molecule or its multimers useful for treating
                                                                                                                                                                                                                                              inflammatory and autoimmune diseases, cancer, rheumatoid arthritis, diabetic retinopathy, obesity, sleep disorders and infertility
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87.0%; Score 20; DB 23; Length 6; 60.0%; Pred. No. 7.8e+05; Ltive 0; Mismatches 2; Indels
                                                                                                                                                      Gudas JM;
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                                                                                                                                                      Boone TC,
                                                                                                                                                                                                                                                                                                           Claim 39; Page 53; 176pp; English.
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                                                                                                                                                      Liu C, Cheetham JC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95WO-GB01719.
                                         02-MAY-2001; 2001WO-US14310.
                                                                             03-MAY-2000; 2000US-0563286
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                                                                                                                                                                                            WPI; 2002-130313/17.
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Matches 3; Conserv
                                                                                                                   (AMGE-) AMGEN INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          6 AA;
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 08-NOV-2001.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence
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                                                                                                                                                        Feige U,
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New peptides of the invention which interfere with the binding of the p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein binding motif between amino acids 16-33 (AAR89914). Peptides contg. these sequence were identified by immobilising a library of peptides amino acids long) covering the mouse or human p53 protein and adding an extract contg. expressed MDM2 protein. The interaction was identified by the antibody D0-1. This is the sequence of a human p53 epitope for the antibody. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Interference; binding; p53; tumour suppressor; MDM2; oncogene; cancer; inhibition; diagnosis; treatment; malignancy; consensus; antibody.
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                                                                                                                                                                                                                                              New cpds, which interfere with binding of MDM2 and p53 - used to develop prods. for use in the diagnosis and treatment of cancer and
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 7;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Indels
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60.0%; Pred. No. 7.8e+05;
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                                                                                                                                                                                                                                                                                                                                                                  Example 3; Page 26; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR89925 standard; peptide; 7 AA.
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                          94US-0277660.
95US-0424957.
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les 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Lane DP, Picksley SM;
                                                                                                                                        Lane DP, Picksley SM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WPI; 1996-105905/11.
                                                                                (UYDU-) UNIV DUNDEE.
                                                                                                                                                                                              WPI; 1996-105905/11.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (UYDU-) UNIV DUNDEE
                                                                                                                                                                                                                                                                                                            other malignancies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 19-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20-JUL-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20-JUL-1994;
19-APR-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 FSDLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WO9602642-A1
                          20-JUL-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-FEB-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AAR89925;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Matches
                RESULT 11
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The invention relates to a synthetic fibronectin type III (Fn3) bolypeptide monobody that comprises Fn3 beta-strand domain sequences that are linked to loop region sequences (LRSS). One or more of the loop sequences in the synthetic Fn3 vary by deletion, insertion, or replacement of at least 2 amino acids from the corresponding LRSS in wild-type Fn3. Host cells containing an expression vector comprising the synthetic Fn3 nucleic acid are used for the production of the Fn3 monobody. The invention also provides methods of identifying the amino acid sequence of a polypeptide molecule (i) capable of binding to a specific binding partner (SBP) so as to form a polypeptide SBP complex, (ii) capable of catalysing a chemical reaction with a catalysed rate constant, Kcat, and an uncatalysed rate constant, Kucat, and an uncatalysed rate constant, Kucat, such that the ratio of the Kcat/Kucat is greater than 10. Sequences AAM94139-46 represent FG loop sequences of yeast ubiquitin-binding monobodies.
New peptides of the invention which interfere with the binding of the p53 tumour suppressor protein and the MDM2 oncogene protein contain the peptide sequences AAR89911-3 which form part of the p53 protein binding motif between amino acids 16-33 (AAR89914). Peptides contg. these sequence were identified by immobilising a library of peptides amino acids long) covering the mouse or human p53 protein and adding an extract contg. expressed MDM2 protein. This sequence represent the key amino acids in the p53 binding site required for mouse MDM2 binding. The peptides and methods of identifying similar inhibitory peptides can be used to diagnose and treat e.g. cancer and other malignancies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Fibronectin type III; Fn3; monobody; beta-strand domain; loop region; specific binding partner; SBP; catalysis; LRS; ubiquitin.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Production of antibody compounds, particularly catalytic antibodies - using a fibronectin type III molecular scaffolding comprising beta-strand domain sequences and modified in one or more loop
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87.0%; Score 20; DB 17; Length 7; 60.0%; Pred. No. 7.8e+05; ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FG loop sequence of ubiquitin-binding monobody clone 213.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (RESE ) RESEARCH CORP TECHNOLOGIES INC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Example 10; Page 41; 96pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW94141 standard; peptide; 7 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97US-0049410.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-APR-1999 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Saccharomyces cerevisiae.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 1999-060331/05.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sest Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3 FSDLW 7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             12-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAW94141;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          sednences
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Koide S;
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                                                                                                                                                                                                                                                                                                                                                                                                         Transcriptional adapter motif; TRAM; TRAM interaction motif; TRIM; creb binding protein; transcriptional regulation; cytostatic; antiviral; cell cycle inhibitor; viral transcription inhibitor; cancer; tumour; viral disease; viral infection; cell cycle; apoptosis; growth arrest.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    useful for preparing alpharaceutical composition and for disrupting an interaction between TRAM sequence and TRIM sequence in vitro, thereby inhibiting viral transcription or call cycle progression in mammalian cells especially cancer cell. Compounds which disrupt interaction between TRIMTRAM containing polypeptides can be used therapeutically to prevent or treat viral diseases and tumours. The polypeptides reduce susceptibility of cells to viral infection and regulate cell cycle antibodies against the TRIM or TRAM sequences. HPV types associated with high risk or low risk of cervical cancer can be distinguished with high risk or low risk of cervical cancer can be distinguished (CBP) TRAM sequence. The present sequence represents a TRIM sequence given in the present invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  The present invention describes a polypeptide comprising a transcriptional adaptor motif (TRAM) or a TRAM-interaction motif (TRAM) or a TRAM-interaction motif (TRAM) (1). (2) can be used for identifying compounds (11) comprising a polypeptide capable of disrupting an interaction between a TRAM sequence and/or a TRAM sequence Polypeptides which bind to a TRAM sequence or a TRAM sequence are identified by incubating, the polypeptide with (1) and determining if the polypeptide interacts with (1). Compound (11) is
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel polypeptides cells useful for treating viral disease and cancer
                                                                                .0
                                           Length 7;
                                     87.0%; Score 20; DB 20; Length 7; 60.0%; Pred. No. 7.8e+05; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                        TRAM-interaction motif derived from p53 SEQ ID NO:12.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MOLE-) INST MOLECULAR & CELL BIOLOGY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Disclosure; Page 7; 73pp; English.
                                                                                                                                                                                                                                                     AAY57789 standard; peptide; 7 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                99WO-GB01668.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99GB-0000157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   98GB-0011303
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Zimmermann H;
                                                                                                                                                                                                                                                                                                                               20-MAR-2000 (first entry)
                                                       Local Similarity 60.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     WPI; 2000-072620/06.
7 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                O'Connor MJ,
                                                                                                                   1 FXXLW 5
                                                                                                                                                        1 FADLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Unidentified.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               WO9961608-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            26-MAY-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     05-JAN-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       02-DEC-1999
  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                         Query Match
                                                                              Matches
                                                                                                                                                                                                                RESULT 12
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Score 20; DB 21; Length 7; Pred. No. 7.8e+05;

87.0%;

Query Match Best Local Similarity

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WO200121770-A2
                                                                                                                                                                                       Mushahwar IK,
                                                                                                              29-JUL-1994;
23-NOV-1994;
23-NOV-1994;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      11-JUN-2001
                                                                             07-JUN-1995;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Glycine max.
                                                                                              14-FEB-1994;
13-MAY-1994;
                                            US6051374-A.
                                                             18-APR-2000
                                                                                                                                         30-JAN-1995
                                                                                                                                                                                                                                                    target HGBV
                                                                                                                                                                              Dawson GJ,
                                                                                                                                                                                                                                                                                                                                                                                                 invention
                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAB62220;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 15
AAB62220
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                     The present invention relates to fibronectin type III (Fn3) molecules comprising a stabilising mutation as compared to a wild-type Fn3. Fn3 can be used as a scaffold to engineer artificial binding proteins. Modifications of the Fn3 scaffold that increase its stability are useful in that they allow the introduction of more mutations for better functions, and that these make it possible to use Fn3-based engineered proteins in a wider range of applications. The present sequence is a peptide described in the exemplification of the invention.
ó
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
Gaps
                                                                                                                                                                                                                                                                                                                                  New fibronectin type III molecule comprising a stabilizing mutation, useful for introducing more mutations for better functions, and in a wider range of applications
                                                                                                                                                   Fibronectin type 3; mutant; stabilising mutation; Fn3; antibody;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         .
ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 20; DB 23; Length 7; Pred. No. 7.8e+05; Indels 0; Mismatches 2; Indels
Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Hepatitis GB virus protein sequence SEQ ID NO:252.
2;
Mismatches
                                                                                                                                                                                                                                                                      (RESE ) RESEARCH CORP TECHNOLOGIES INC. (KOID/) KOIDE S.
                                                                                                                                   Ubiquitin binding monobody 213 FG loop.
                                                                                                                                                                                                                                                                                                                                                                      Example 10; Page 138; 164pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB09130 standard; Protein; 8 AA.
                                                                               ABB94392 standard; Peptide; 7 AA.
ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87.0%;
                                                                                                                                                                                                                                    11-JUL-2001; 2001WO-US21855
                                                                                                                                                                                                                                                     11-JUL-2000; 2000US-217474P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-AUG-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
 Conservative
                                                                                                                                                                                                                                                                                                                 WPI; 2002-171708/22.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match .
Best Local Similarity
Matches 3; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 AA;
                                                                                                                                                              binding protein.
                                                                                                                                                                                                 WO200204523-A2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 FXXLW 5
                  FXXLW 5
                                   S
                                                                                                                                                                              Unidentified
                                                                                                                 12-JUN-2002
                                                                                                                                                                                                                  17-JAN-2002
3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 FADLW
                                   FSDLW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           AAB09130;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                 ABB94392;
                                                                                                                                                                                                                                                                                                 Koide S;
Matches *
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 14
                                                             RESULT 13
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                                                                       ABB94392
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The present invention describe a method for detecting target hepatitis GB virus (HGBV) nucleic acid (THN) in a test sample (T) subspected of contraining HGBV. The method involves reacting (T) with a HGBV polynucleotide probe (I) containing 15 contiguous nucleotides, and which selectively hybridises to the HGBV genome or its full complement, and detecting the complex that contains THN, indicating the presence of target HGBV. The method is used for detecting target HGBV nucleic acid in the test sample suspected of containing HGBV and for characterisation of newly ascertained etiological agent of non-A, non-B, non-C, non-D and non-B hapatitis causing agents collectively termed as hepatitis GB virus. AAA55270 to AAA55489 and AAB089885 to AAB09480 represent nucleotide and protein sequences used in the exemplification of the present
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Soybean, glutathione-S-transferase, GST, homoglutathione synthetase, HGS, beta-alanine, gamma-glutamylcysteine, herbicide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Detecting target hepatitis GB virus nucleic acid in a test sample suspected of containing HGBV comprises reacting the test sample the HGBV polynucleotide probe and detecting the complex that contains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ;
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Pilot-Matias TJ, Buijk S
Erker JC, Schlauder GG;
Hepatitis GB virus; HGBV; diagnosis; therapeutic; immunogenic;
infection; detection; characterisation; hepatitis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      87.0%; Score 20; DB 21; Length 8; 60.0%; Pred. No. 7.8e+05; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Glycine max homoglutathione synthetase protein region.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Example 9; Column 331-332; 369pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Muerhoff AS,
JN, Desai SM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                AAB62220 standard; peptide; 8 AA.
                                                                                                                                                                                                                                                                                                                                                                                      94US-0196030.
94US-0242654.
94US-0283314.
94US-0344185.
94US-0344190.
95US-0377557.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Simons JN,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Leary TP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 2000-338307/29.
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                                                                                                        Hepatitis GB virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (ABBO ) ABBOTT LAB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          8 AA;
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29-MAR-2001.
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18-SEP-2000; 2000WO-GB03573.

99GB-0022346. 21-SEP-1999;

(ZENE) ZENECA LTD.

Townson JK, Edwards R, Cummins I, Skipsey M; Andrews CJ, Jepson I,

WPI; 2001-257978/26.

sequences from soybean for producing plants which are resistant and tolerant to herbicide comprising fomesafen and/or acifluorfen Novel glutathione-S-transferase and homoglutathione synthetase

Claim 9; Page 39; 64pp; English.

The invention relates to new soybean glutathione-S-transferase (GST) and a homoglutathione synthetase (HGS). The HGS is capable of catalyzing the addition of beta-alanine onto gamma-glutamylcysteine. Polynucleotides encoding the proteins of the invention are useful for producing plants which are resistant and/or tolerant to a herbicide comprising fomesafen plants with further desired agronomic trait, especially resistant to a herbicide, comprising glyphosate or its salt. Further desired agronomic traits altered field, altered nutritional value, altered quality or any other desirable agronomic trait. GST or its variant is also useful as a selectable marker gene. The present sequence represents a Glycine max HGS protein fragment.

8 AA; Sequence

0; Gaps Query Match 87.0%; Score 20; DB 22; Length 8; Best Local Similarity 60.0%; Pred. No. 7.8e+05; Matches 3; Conservative 0; Mismatches 2; Indels

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1 FXXLW 5 δ 2 FAGLW 6 g Search completed: February 12, 2003, 11:13:44 Job time : 13.5 secs

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Sequence 83, Appli
Sequence 5, Appli
Sequence 1, Appli
Sequence 36, Appl
Sequence 36, Appl
Sequence 252, App
Sequence 11, Appl
Sequence 14, Appl
Sequence 37, Appl
Sequence 38, Appl
Sequence 38, Appl
Sequence 38, Appl
Sequence 38, Appl
Sequence 36, Appl
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Appli
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                                                                                                 February 12, 2003, 11:15:02 ; Search time 3.33333 Seconds (without alignments) 38.323 Million cell updates/sec
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Sequence 6
Sequence 6
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1: /cgn2_6/ptodata/1/pubpaa/USO8_NEW_PUB.pep:*
2: /cgn2_6/ptodata/1/pubpaa/PUT_NWW_PUB.pep:*
3: /cgn2_6/ptodata/1/pubpaa/PUT_NWW_PUB.pep:*
4: /cgn2_6/ptodata/1/pubpaa/USO6_NEW_PUB.pep:*
5: /cgn2_6/ptodata/1/pubpaa/USO7_NEW_PUB.pep:*
6: /cgn2_6/ptodata/1/pubpaa/USO7_PUBCOMB.pep:*
7: /cgn2_6/ptodata/1/pubpaa/USO7_PUBCOMB.pep:*
8: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
9: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
10: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
11: /cgn2_6/ptodata/1/pubpaa/USO9_PUBCOMB.pep:*
13: /cgn2_6/ptodata/1/pubpaa/USO0_PUBCOMB.pep:*
14: /cgn2_6/ptodata/1/pubpaa/USO0_PUBCOMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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0 US-09-732-384-4

0 US-09-732-384-5

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US-09-903-412-36

0 US-09-905-749A-36

0 US-09-844-456-8

US-09-844-456-11

US-09-844-456-11

US-09-844-456-11

US-09-844-456-11

US-09-844-456-11

US-09-214-371-38

0 US-09-214-371-35

0 US-09-214-371-37

US-09-214-371-17

US-09-214-371-17
                                                                                                                                                                                                                                                                                                                         Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                          140259 segs, 25548876 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUMMARIES
                                                                                                                                                                                                                                                                                                                                                                                                                Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                  OM protein - protein search, using sw model
                                                                                                                                                                                                                                        BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                     US-09-403-440A-1
23
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Match Length DB
                                                                                                                                                                                                        1 FXXLW 5
                                                                                                                                                                                                                                                                                                                                                             Minimum DB seq length: 0 Maximum DB seq length: 25
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                                                                                                                                                                                                           Sequence:
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Sequence 203,
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US-09-214-371-83
                                                                                Sequence
Sequence
Sequence
Sequence
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Sequence
Sequence
Sequence
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0
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US-09-214-371-39
US-09-214-371-40
US-09-214-371-40
US-09-214-371-14
US-09-214-371-14
US-09-732-384-7
US-10-155-059-18
US-10-155-059-10
US-10-155-059-14
US-10-155-059-14
US-10-155-059-15
US-10-155-059-15
US-10-155-059-16
                                                                                                                                                     US-09-774-639-203

US-09-214-371-2

US-09-214-371-3

US-09-81-77-3

US-09-862-260A-9

US-09-81-276-5

US-09-81-276-5
                                                                                                                                                                                                                                                                                                                            APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Nagelica
APPLICANT: Bottger, Angelica
APPLICANT: Picksley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTON: Inhibitors of the Interac
CURRENT APPLICATION NUMBER: US/09/214,3718
CURRENT FILING DATE: 1999-03-26
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
SENGTH: 6
                                                                                                                                                                                                                                                         ALIGNMENTS
                                                                                                                                                                                                                                                                                                    ; Sequence 83, Application US/09214371B
; Patent No. US20010018511A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: PRT
ORGANISM: Artificial Sequence
  3; Conservative
   Query Match
Best Local Similarity
Matches 3; Conserval
 US-09-214-371-83
  FEATURE:
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; Sequence 4, Application US/09732384

RESULT 2 US-09-732-384-4

1 FXXLW | | |

8 6

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Matches
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APPLICANT: Dost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: MBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
ANTIBODIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ..
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                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: Description of Artificial Sequence:Protein OTHER INFORMATION: fragment not in inhibitory p53 polypeptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

87.0%; Score 20; DB 10; Length 6;
Best Local Similarity 60.0%; Pred. No. 1.2e+05;
Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                h 87.0%; Score 20; DB 10; Length 6; Similarity 60.0%; Pred. No. 1.2e+05; 3; Conservative 0; Mismatches 2; Indels
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US-09-132-384-5
i Sequence 5, Application US/09732384
j Patent No. US20020132977A1
j GENERAL INFORMATION:
i APPLICANT: Van, Zhi-Min
j APPLICANT: Gu, JiJie
j TITLE OF INVENTION: Inhibition of p53 Degradation
j FILE REFERENCE: 21508-044
j CURRENT APLICATION NUMBER: US/09/732,384
j CURRENT APPLICATION NUMBER: 60/169,816
pRIOR APPLICATION NUMBER: 60/169,816
pRIOR FILING DATE: 1999-12-08
j NUMBER OF SEQ ID NOS: 10
j SOFTWARE: PatentIn Ver. 2.1
                GENERAL INFORMATION:
APPLICANT: Yuan, Zhi Min
APPLICANT: Yuan, Zhi Min
TITLE OF INVENTION: Inhibition of p53 Degradation
FILE REFERENCE: 21508-044
CURRENT APPLICATION NUMBER: US/09/732,384
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 60/169,816
PRIOR PILING DATE: 1999-12-08
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver: 2.1
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 1, Application US/10155059
Patent No. US20020147173A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
ORGANISM: Artificial Sequence
Patent No. US20020132977A1
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Best Local Similarity
Matches 3; Conserv
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                                                                                                                                                                                                                               COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/155,059
FILING DATE: 24 May-2002
CLASSIFICATION: <UNKnown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Koide, Shohei
TITLE OF INVENTION: ARTIFICIAL ANTIBODY FOLYPEPTIDES
FILE REFERENCE: 109.050US1
CURRENT APPLICATION NUMBER: US/09/903,412
CURRENT FILING DATE: 2001-07-11
PRIOR APPLICATION NUMBER: US 60/217,474
PRIOR PILING DATE: 2000-07-11
SOFTWARE: FASTSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                      CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-345-6054
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 36, Application US/09903412; Publication No. US20030027319A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEX: <URANDOMS INFORMATION FOR SEQ ID NO: 1: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT
ORGANISM: Artificial Sequence
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3; Conservative
                                                                                                                                         USA
                                                                                            CITY: Boston
                                                                                                                                                                ZIP: 02110
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Best Local Similarity
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Best Local Similarity
                                                                                                              STATE: MA
COUNTRY: U
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US-09-903-412-36
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APPLICANT: JAMES C. ERKER
APPLICANT: STREI L. BULJK
APPLICANT: STREI L. BULJK
APPLICANT: STREI L. BULJK
TITLE OF INVENTION: NON-A, NON-B. NON-C, NON-D, NON-E HEPATITIS
TITLE OF INVENTION: REAGENTS AND METHODS FOR THEIR USE
CORRESPONDENCES ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87.0%; Score 20; DB 8; Length 8; 60.0%; Pred. No. 1.2e+05; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                            COUNTRY: USA

COUNTRY: USA

COMPUTER REDABLE FORM:
MEDIUT TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,550B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Houghton, Michael
APPLICANT: Houghton, Michael
APPLICANT: Choo, Oui-Lim
APPLICANT: Choo, Oui-Lim
APPLICANT: Rio, George
TITLE OF INVENTION: Hopatitis C virus protease
FILE REFERENCE: 223002010005
CURRENT APPLICATION NUMBER: US/09/884,456
CURRENT FILING DATE: 1999-02-19
PRIOR APPLICATION NUMBER: 08/709,177
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR PRIUNG DATE: 1995-05-12
PRIOR PELING DATE: 1991-04-04
PRIOR PELING DATE: 1991-04-04
PRIOR APPLICATION NUMBER: 07/680,296
PRIOR APPLICATION NUMBER: 07/680,296
PRIOR APPLICATION NUMBER: 07/680,433
                                                                                                                                                                                     ADDRESSEE: ABBOTT LABORATORIES D377/AP6D STREET: 100 ABBOTT PARK ROAD CITY: ABBOTT PARK STATE: IL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR FILING DATE: 1990-04-04
NUMBER OF SEQ ID NOS: 90
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 435435
ATTORNEY/AGENT INFORMATION:
NAME: POREMBSKI, RRISCILLA E.
REGISTRATION NUMBER: 33,207
REFERENCE/DOCKET NUMBER: 5527.PC.01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 708-937-6365
             ANTHONY SCOTT MUERHOFF
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 8, Application US/09884456; Publication No. US20030027317A1; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 60.0'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 708-938-2623
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 8 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: 708-22623
TELEPHONE: 708-938-2623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: protein US-08-424-5508-252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                amino acid
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87.0%; Score 20; DB 10; Length 7;

Best Local Similarity 60.0%; Pred. No. 1.2e+05;

Matches 3; Conservative 0; Mismatches 2; Indels
           2; Indels
                                                                                                                                                                                                      Sequence 36, Application US/09096749A
Patent No. US20020019517A1
GENERAL TREOMATION:
APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
ANUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A.
STREET: 121 South Eighth Street, Ste. 1600
         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Ann S. Vikenins
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US1
TELECOMMUNICATION INFORMATION:
TELEPHONE: (612) 373-6900
TELEPHONE: (612) 339-3061
INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 252, Application US/08424550B
Fatent No. US20020119447A1
GENERAL INFORMATION:
APPLICANT: JOHN N. SIMONS
APPLICANT: TAMI J. PILOT-MATIAS
APPLICANT: GEORGE J. DAWSON
APPLICANT: GEORGE G. SCHLAUDER
APPLICANT: SURBSH M. DESAI
APPLICANT: THOMAS P. LEARY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/096,749A FILING DATE: June 12, 1998
PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
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Matches ' 3; Conservative
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ZIP: 55402
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ANTI-SENSE: NO
; FRAGMENT TYPE: 1
; ORIGINAL SOURCE:
US-09-096-749A-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-08-424-550B-252
                                                   1 PXXLW 5
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                                                                                             1 FADLW 5
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87.0%; Score 20; DB 9; Length 9;
Best Local Similarity 60.0%; Pred. No. 1.2e+05;
Matches 3; Conservative 0; Mismatches 2; Indels
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Best Local Similarity 60.0%; Pred. No. 1.2e+05;
Matches 3; Conservative 0; Mismatches 2; Indels
                                              TYPE: PRT
ORGANISM: West Nile Fever virus
FEATURE:
OTHER INFORMATION: West Nile Fever virus protease
US-09-884-456-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Houghton, Michael
APPLICANT: Choo, Oui-Lim
APPLICANT: Choo, Oui-Lim
TITLE GOT INVENTION: Hopetitis C virus protease
FILE REFERENCE: 223002010005
CURRENT APPLICATION NUMBER: US/09/884,456
CURRENT FILING DATE: 1990-02-19
PRIOR APPLICATION NUMBER: 08/709,177
PRIOR APPLICATION NUMBER: 08/709,177
PRIOR FILING DATE: 1996-09-06
PRIOR APPLICATION NUMBER: 08/40,548
PRIOR PRILING DATE: 1991-04-04
PRIOR PILING DATE: 1991-04-04
PRIOR PILING DATE: 1991-04-04
PRIOR FILING DATE: 1991-04-04
PRIOR FILING DATE: 1990-04-04
PRIOR FILING DATE: 1900-04-04
PRIOR PILING DATE: 1900-04-04
PRIOR FILING DATE: 1900-04-04
PRIOR PILING DATE: 1900-04-04
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; OTHER INFORMATION: Murray Valley virus protease
US-09-884-456-11
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CURRENT APPLICATION NUMBER: US/09/884,456
PRIOR APPLICATION NUMBER: 09/253,230
                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 11, Application US/09884456; Publication No. US20030027317A1; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Murray Valley virus
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APPLICANT: Choo, Oui-Lim
APPLICANT: Kuo, George
                                                                                                                                                                                                                                                                                              1 FXXLW 5
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US-09-884-456-11
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US-09-884-456-14
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; SEQ ID NO 8
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87.0%; Score 20; DB 10; Length 9;
Best Local Similarity 60.0%; Pred. No. 1.2e+05;
Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 87.0%; Score 20; DB 9; Length 9; Best Local Similarity 60.0%; Pred. No. 1.2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
PRIOR FILING DATE: 1999-02-19
PRIOR APPLICATION NUMBER: 08/709,177
PRIOR FILING DATE: 1996-09-06
PRIOR APPLICATION NUMBER: 08/440,548
PRIOR PILING DATE: 1995-05-12
PRIOR PILING DATE: 1995-05-12
PRIOR PELING DATE: 1994-12-06
PRIOR FILING DATE: 1991-04-04
PRIOR FILING DATE: 1991-04-04
PRIOR PELING DATE: 1991-04-04
PRIOR PELING DATE: 1991-04-04
PRIOR FILING DATE: 1990-04-04
PRIOR FILING DATE: 1990-04-04
PRIOR FILING DATE: 1990-04-04
PRIOR FILING DATE: 1990-04-04
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CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Kunjin virus protease
US-09-884-456-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Pickeley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Hochkeppel, Heinz-Kurt
APPLICANT: Garcia-Echeverria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 37, Application US/09214371B ; Patent No. US20010018511A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: X = Ac=Cys (Acrd)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Lane, David
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                       TYPE: PRT
ORGANISM: Kunjin virus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME/KEY: VARIANT LOCATION: (9)
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LENGTH: 9
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APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Hockeppel, Heinz-Kurt
APPLICANT: Hockeppel, Heinz-Kurt
APPLICANT: Garcia-Cheverria, Carlos
APPLICANT: Furet, Paecal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REFERENCE: 4-20337/A/PCT
CURRENT APPLICATION NUMBER: PCT/S99/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR APPLICATION NUMBER: PCT/EP97/03549
NUMBER OF SEQ ID NOS: 83
SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 35
LENGTH: 10
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                                                                                                                                                         APPLICANT: DOUGSEY, ADSTITUTE APPLICANT: PLOKESEY, Stephen APPLICANT: Chene, Patrick APPLICANT: Chene, Patrick APPLICANT: Chene, Patrick APPLICANT: Garcia-Echeverria, Carlos APPLICANT: Garcia-Echeverria, Carlos APPLICANT: Furet, Pascal Inhibitors of the Interaction of P53 and MDM2 TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2 CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
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PEATURE:
OTHER INFORMATION: Description of Artificial Sequence:peptide
NAME/KEY: VARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Description of Artificial Sequence:peptide NAME/KEY: VARIANT LOCATION: (1)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 87.0%; Score 20; DB 10; Length 9; Best Local Similarity 60.0%; Pred. No. 1.2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                     Bottger, Volker
Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
Hochkeppel, Heinz-Kurt
Gardia-Echeverria, Carlos
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                      Sequence 38, Application US/09214371B Patent No. US20010018511A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; LOCATION: (1)
; OTHER INFORMATION: x = Ac-Cys
; NAME/KEY: VARIANT
; LOCATION: (9)
; OTHER INFORMATION: X = Pro-NH2
US-09-214-371-38
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Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 38
LENGTH: 9
                                                              GENERAL INFORMATION:
APPLICANT: Lane, D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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US-09-214-371-38
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Gaps
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CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT APPLICATION NUMBER: US/09/214,371B
CURRENT FILING DATE: 1999-03-26
PRIOR APPLICATION NUMBER: PCT/EP97/03549
PRIOR FILING DATE: 1997-07-04
NUMBER OF SEQ ID NOS: 83
SSCTWARE: Patentin Ver. 2.0
LENGTH: 10
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                                                                                                                                   87.0%; Score 20; DB 10; Length 10; 60.0%; Pred. No. 1.4e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87.0%; Score 20; DB 10; Length 10; 60.0%; Pred. No. 1.4e+02; tive 0; Mismatches 2; Indels
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Bottger, Volker
Bottger, Angelica
Picksley, Stephen
Chene, Patrick
Hochkeppel, Heinz-Kurt
Garcia-Echeverria, Carlos
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Hochkeppel, Heinz-Kurt
Garcia-Echeverria, Carlos
Furet, Pascal
                                                                                                                                                                                                                                                                                                                                                    ; Sequence 36, Application US/09214371B ; Patent No. US20010018511A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 17, Application US/09214371B
; Patent No. US20010018511A1
LOCATION: (1)
CTHER INFORMATION: X = Ac-Cys(Acrd)
NAME/KEX: VARIANT
LOCATION: (10)
COTHER INFORMATION: X = Pro-NH2
US-09-214-371-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; LOCATION: (10); OTHER INFORMATION: x = Pro-NH2 US-09-214-371-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME/KEY: VARIANT
LOCATION: (1)
OTHER INFORMATION: X = AC-CYS
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Bottger, Angelica
Picksley, Stephen
Chene, Patrick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: PRT ORGANISM: Artificial Sequence
                                                                                                                                     Query Match 87.0
Best Local Similarity 60.0
Matches 3; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Furet, Pascal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Lane, David
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
nes 3; Conserv
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APPLICANT: Lane, Da
APPLICANT: Bottger,
APPLICANT: Pottger,
APPLICANT: Conse,
                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION
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                                                                                                                                                                                                                  1 FXXLW 5
                                                                                                                                                                                                                                                         5 FSDLW 9
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APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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FILE REFERENCE: 4-20037/APCT
CURRENT APPLICATION NUMBER: 0.5/09/214,371B
CURRENT APPLICATION NUMBER: 0.5/09/214,371B
CURRENT FILING DATE: 1999-03-26
FRIOR APPLICATION NUMBER: PCT/EP97/03549
FRIOR APPLICATION NUMBER: PCT/EP97/03549
NUMBER OF SEQ ID NOS: 83
SOFTWARE: PatentIn Ver. 2.0
LENGTH: 12
TYPE: PRT
CRAMMER: PATENTIAL SEQUENCE
FEATURE: ORGANISM: Artificial Sequence
FEATURE:
CORPER INFORMATION: Description of Artificial Sequence:peptide
WS-09-214-371-17
QUARTY MATCH
SEX INFORMATION: Description of Artificial Sequence:peptide
WATCHES 3; CONSERVATION: 0; Mismatches 2; Indels 0; Gaps

QY 1 FXXLW 5
Db 4 FSDLW 8
Search completed: February 12, 2003, 11:25:07
Job time: 4.33333 secs
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Sequence 16, Appl
Sequence 11, Appl
Sequence 11, Appl
Sequence 114, App
Sequence 11, Appl
Sequence 11, Appl
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                                                                                                                                                February 12, 2003, 11:13:52 ; Search time 99.375 Seconds (without alignments) 32.439 Willion cell updates/sec
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1: /cgn2_6/ptodata/1/paa/PCTUS_COMB.pep:*
2: /cgn2_6/ptodata/1/paa/USOG_COMB.pep:*
4: /cgn2_6/ptodata/1/paa/USOG_COMB.pep:*
5: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
6: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
7: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
7: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
8: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
9: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
10: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
11: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
12: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
13: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
14: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
15: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
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16: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
17: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
18: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
19: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
20: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
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23: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
24: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
25: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
26: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
27: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
28: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
29: /cgn2_6/ptodata/1/paa/USOB_COMB.pep:*
GenCore version 5.1.3
(c) 1993 - 2003 Compugen Ltd.
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US-09-936-956-10

PCT-US99-02405-11

PCT-US99-26277-114

US-09-601-558-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                     4569144 segs, 644733110 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                     - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                        BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                     US-09-403-440A-1
23
1 FXXLW 5
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Match Length
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Minimum DB seq length: 0
Maximum DB seq length: 25
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91.3
87.0
87.0
87.0
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                                                                                                  OM protein
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                                                                                                                                                                                                                                                                                                        Sequence:
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Sequence 114, App
Sequence 11, Appl
Sequence 2, Appl
Sequence 3, Appli
Sequence 83, Appli
Sequence 83, Appli
Sequence 130, App
                                                                                                                                           Sequence 5, Appli
Sequence 1, Appli
Sequence 27, Appl
Sequence 36, Appl
Sequence 36, Appl
Sequence 24, Appl
                                                                                                                                                                                                                                                                     Sequence 206, App
Sequence 604, App
Sequence 381, App
Sequence 123, App
Sequence 123, App
Sequence 123, App
Sequence 252, App
                                                                                                                                                                                                                                                               App
                                                                                                                                Sequence 4, Appl
                                                                                                                                                                                                                                Sequence 36, A Sequence 36, A Sequence 304, A Sequence 206, A Sequence 293, A Sequence 694, A Sequence 621, A
                                                                                           Sequence 130,
Sequence 130,
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Sequence
Sequence
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US-08-277-660-16

Sequence 16, Application US/08277660

GENERAL INFORMATION:

APPLICANT: Dicksley, Steven M.

APPLICANT: Lane, David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof

NUMBER OF SEQUENCES:

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400 STREET: Four Embarcadero Center, Suite 3400 STATE: San Francisco STATE: California COUNTRY: United States ZIP: 94111-4187 COMPUTER READSHE FORM: MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660
FILING DATE: 20-JUL-1994
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
2 US-09-831-047C-114
2 US-09-894-594-11
US-08-277-660-3
6 US-08-277-660-3
6 US-08-277-660-3
6 US-09-276-60-3
6 US-09-428-082-130
9 US-09-53-868-130
9 US-09-53-868-130
9 US-09-53-868-130
1 US-09-732-384-5
2 US-09-732-384-5
2 US-09-732-384-5
2 US-09-732-384-5
2 US-09-732-384-5
2 US-09-732-384-5
2 US-09-732-382-7
2 US-09-762-7660-2
2 US-09-762-7660-2
2 US-09-762-7660-2
2 US-08-344-1856-123
2 US-08-344-1856-123
2 US-08-344-1856-123
2 US-08-344-1856-123
2 US-08-446-749-252
2 US-08-486-749-252
2 US-08-486-749-252
2 US-08-486-746-252
2 US-08-486-746-252
2 US-08-486-746-252
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US-09-458-297-345
US-09-458-297-475
US-09-458-297-690
US-09-458-297-690
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REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-60244/WHD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Trustees of the University of Pennsylvania TITLE OF INVENTION: Peptide Mimotopes of Carbohydrate Antigens FILE REPRENCE: upn3544 CURRENT APPLICATION NUMBER: PCT/US99/02405 CURRENT FILING DATE: 1999-02-04 EARLIER APPLICATION NUMBER: 60/073,690 EARLIER, FILING DATE: 1998-02-04 NUMBER OF SEQ ID NOS: 60 SOFTWARE; Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: KuDOS Pharmaceuticals Limited
APPLICANT: KuDOS Pharmaceuticals Limited
APPLICANT: Jackson, Stephen P
APPLICANT: Jackson, Stephen P
TITLE OF INVENTION: Screening Methods Based on FHA Domains
FILE REFERENCE: MEWBOLK
CURRENT APPLICATION NUMBER: US/09/936,956
CURRENT APPLICATION NUMBER: GB 9906432.1
PRIOR FILING DATE: 1999-00-19
PRIOR FILING DATE: 1999-06-28
PRIOR FILING DATE: 1999-06-28
NUMBER OF SEQ ID NOS: 57
SOFTWARE: PATENTIN VET: 2.1
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91.3%; Score 21; DB 23; Length 15;
Best Local Similarity 60.0%; Pred. No. 7.7e+03;
Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                  Query Match
91.3%; Score 21; DB 6; Length 11;
Best Local Similarity 60.0%; Pred. No. 6.1e+03;
Matches 3; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 10, Application US/09936956 GENERAL INFORMATION:
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US-09-936-956-10
                TELEPHONE: (415) 781-1989
TELEX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acids
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ORGANISM: Artificial Sequence
                                                                                                                                                                                                                          TOPOLOGY: linear
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LENGTH: 5 ,
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0; Gaps
                                                   ; OTHER INFORMATION: Description of Artificial Sequence: Novel Sequence PCT-US99-02405-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: The Mistar Institute of Anatomy and Biology
APPLICANT: The Mistar Institute of Anatomy and Biology
APPLICANT: The Trustees of the University of Pennsylvania
APPLICANT: Blaszczyk-Thurin, Maddalena
APPLICANT: Keiber-Emmons, Thomas
TITLE OF INVENTION: Compositions and Methods For Treatment of CancerFILE REPERBNCE: WST93PCT
CURRENT APPLICATION NUMBER: PCT/US99/26277
CURRENT FILING DATE: 1999-11-05
EARLIER FILING DATE: 1999-11-06
SARLIER FILING DATE: 1999-11-06
NUMBER OF SEQ ID NOS: 12.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Kieber-Emmons, Thomas
APPLICANT: Kieber-Emmons, Thomas
APPLICANT: The Trustees of the University of Pennsylvania
TITLE OF INVENTION: Peptide Mimotopes Of Carbohydrate Antigens
TITLE OF INVENTION: Peptide Mimotopes Of Carbohydrate Antigens
FILE REPERENCE: UPN367.
CURRENT APPLICATION NUMBER: US/09/601,558
CURRENT APPLICATION NUMBER: PCT/US99/02405
PRIOR PILING DATE: 1999-02-04
PRIOR FILING DATE: 1999-02-04
NUMBER OF SEQ ID NOS: 60
NUMBER OF SEQ ID NOS: 60
SEQ ID NO 1:
LENGTH: 5
                                                                                                                                                                                                     ;
0
                                                                                                                                     Query Match 87.0%; Score 20; DB 1; Length 5; Best Local Similarity 60.0%; Pred. No. 4.2e+06; Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 87.0%; Score 20; DB 1; Length 5; Best Local Similarity 60.0%; Pred. No. 4.2e+06; Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: CTHER INFORMATION: peptido-mimetic of a Lewis antigen PCT-US99-26277-114
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; Sequence 114, Application PC/TUS9926277; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-601-558-11; Sequence 11, Application US/09601558; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                     1 FXXLW 5
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                           FEATURE:
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Sequence 114, Application US/09831047C

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: The Wister Institute of Anatomy and Biology
APPLICANT: The Trustees of the University of Pennsylvania
APPLICANT: Rieber-Emmons, Thomas
TITLE OF INVENTION: Compositions and Methods For Treatment of Cancer:
FILE REPRENCE: WST93AUSA
CURRENT PILING DATE: 2001-05-03
FRIOR APPLICATION NUMBER: 60/107,478
PRIOR PYLLING DATE: 1998-11-06
NUMBER OF SEQ ID NOS: 121
SOFTWARE: Patentin version 3.1
SOFTWARE: Patentin version 3.1
SEUGH 1996-11-06
LENGTH: SEQ ID NO 114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                    Gaps
                                                                                                                                                                                                                                                                                                   APPLICANT: The Wistar Institute of Anatomy and Biology
APPLICANT: The Trustees of the University of Pennsylvania
APPLICANT: Blasaczyk-Thurin, Magdalena
APPLICANT: Kieber-Emmons, Thomas
TITLE OF INVENTION: Compositions and Methods For Treatment of Cancer FILE REPERENCE: WST93AUSA
CURRENT APPLICATION NUMBER: US/09/831,047B
CURRENT FILING DATE: 2001-08-14
PRIOR APPLICATION NUMBER: 60/107,478
PRIOR APPLICATION NUMBER: 60/107,478
PRIOR PILING DATE: 1998-11-06
NUMBER OF SEQ ID NOS: 12.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 87.0%; Score 20; DB 22; Length 5; Best Local Similarity 60.0%; Pred. No. 4.2e+06; Matches 3; Conservative 0; Mismatches 2; Indels
Score 20; DB 20; Length 5; Pred. No. 4.2e+06; 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CTHER INFORMATION: Description of Artificial Sequence:
CTHER INFORMATION: peptido-mimetic of a Lewis antigen
US-09-831-047B-114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TYPE: PRT
; ORGANISM: peptido-mimetic of a Lewis antigen
US-09-831-047C-114
                                                                                                                                                                                                                                          US-09-831-047B-114; Sequence 114, Application US/09831047B; GENERAL INFORMATION:
  87.0%;
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                                                    Conservative
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Best Local Similarity
  Query Match
Best Local Similarity
Matches 3; Conserv
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GENERAL INFORTION:

JAPLICANT: Kieber-Emmons, Thomas

APPLICANT: Kieber-Emmons, Thomas

APPLICANT: Weiner, David B.

APPLICANT: Weiner, David B.

APPLICANT: Weiner, David B.

JITLE OF INVENTION: Peptide Mimotopes of Carbohydrate Antigens and DNA Molecules Encorning TITLE OF INVENTION: Same

FILE REFERENCE: UPN-3984

CURRENT PAPLICATION NUMBER: US/09/894,594

CURRENT FILING DATE: 2000-10-6-28

PRIOR APPLICATION NUMBER: 60/601,558

PRIOR APPLICATION NUMBER: 60/033,690

PRIOR FILING DATE: 1998-02-04

PRIOR FILING DATE: 1998-02-04

PRIOR FILING DATE: 1998-02-04

PRIOR FILING DATE: 2000-06-28

NUMBER: OF SEQ ID NOS: 70

SEQ ID NOS: 70

SEQ ID NO 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: FEIGE, ULRICH
APPLICANT: CHEETHAM, JANET
APPLICANT: BOONE, THOMAS CHARLES
TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87.0%; Score 20; DB 22; Length 5; 60.0%; Pred. No. 4.2e+06; trive 0; Mismatches 2; Indels
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CURRENT APPLICATION NUMBER: PCT/US99/25044
CURRENT FILING DATE: 1999-10-25
FEARLIER APPLICATION NUMBER: 60/105,371
EARLIER FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 1133
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 130
LENGTH: 6
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RESULT 8
US-09-389-594-11
Sequence 11, Application US/09894594
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc feature
CTHER INFORMATĪON: Novel Sequence
US-09-894-594-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
'-hes 3; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PCT-US99-25044-130
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                                        Sequence 2, Application US/08277660
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Picksley, Steven M.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3, Application US/08277660
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 87.0%; Score 20; DB 6; Length 6; Best Local Similarity 60.0%; Pred. No. 4.2e+06; Matches 3; Conservative 0; Mismatches 2; Indels
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ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660 FILING DATE: 20-JUL-1994 CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Rloppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Malter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-60244/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEPAK: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                              COMPUTER TRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELERAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 2:
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LENGTH: 6 amino acids
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TYPE: amino acid
STRANDEDNESS:
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CITY: San Francisco
GTATE: California
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US-08-277-660-2
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                     US-08-277-660-2
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US-08-277-660-3
RESULT 10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Lane, Volker
APPLICANT: Bottger, Volker
APPLICANT: Bottger, Angelica
APPLICANT: Bottger, Angelica
APPLICANT: Pickeley, Stephen
APPLICANT: Chene, Patrick
APPLICANT: Chene, Patrick
APPLICANT: Garcia-Echeveria, Carlos
APPLICANT: Furet, Pascal
TITLE OF INVENTION: Inhibitors of the Interaction of P53 and MDM2
FILE REPERENCE: 4-20937/APCT
CURRENT PILLING DATE: 1999-03-26
FRIOR APPLICATION NUMBER: US/09/214,371B
CURRENT FILLING DATE: 1997-07-04
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87.0%; Score 20; DB 16; Length 6;
Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2; Indels
                                      FILING DATE: 20-JUL-1994
CLASSIFICATION: 514
ATTONNEY/GRAT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REBERENCE/DOCKET NUMBER: A-60244/WHD
TELECOMMUNICATION INFORMATION:
TELEPAN: (415) 781-1989
TELER: 910 277299
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                   APPLICATION NUMBER: US/08/277,660
FILING DATE: 20-JUL-1994
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-214-371-83; Sequence 83, Application US/09214371B; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 130, Application US/09428082; GENERAL INFORMATION; APPLICANT: LIU CHUAN-FA APPLICANT: FEIGE, ULRICH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: PRT ORGANISM: Artificial Sequence
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 83
LENGTH: 6
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 60.0
les 3; Conservative
                                                                                                                                                                                                                                                                                                                                         amino acid
                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                            STRANDEDNESS:
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Sequence 130, Application US/09428082B

Sequence 130, Application:
APPLICANT: ERIGE, ULRICH
APPLICANT: LIU, CHUAN-FA

TAPLICANT: CHEETHAM, JANET C.
APPLICANT: BOONE, THOMAS CHARLES
TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
TITLE OF INVENTION WHERE: US/09/428,082B
CURRENT FILING DATE: 1999-10-22
PRIOR APPLICATION NUMBER: 60/105,371
PRIOR FILING DATE: 1999-10-23
NUMBER OF SEQ ID NOS: 1133
SOFTWARE: PATENTIN VERSION 3.1

SEQ ID NO 130
; APPLICANT: CHEETHAM, JANET
; APPLICANT: BOONE, THOMAS CHARLES
; TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
; FILER REFERENCE: A-5.7
; CURRENT APPLICATION NUMBER: US/09/428,082
; CURRENT PILING DATE: 1999-10-22
; EARLIER APPLICATION NUMBER: 60/105,371
; EARLIER PILING DATE: 1998-10-23
; NUMBER OF SEQ ID NOS: 1133
; SOFTWARE: PALENTIN VEY: 2.1
; SEQ ID NO 130
; LENGTH: 6
                                                                                                                                                                                                                                                                                                                                                                                                                      CTHER INFORMATION: Description of Artificial Sequence: MDM/HDM; OTHER INFORMATION: ANTAGONIST PEPTIDE US-09-428-082-130
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APPLICANT: LIU, CHTAN-FA
APPLICANT: CHESTHAM, JANET C.
APPLICANT: CHESTHAM, JANET C.
APPLICANT: BCONE, THOMAS CHARLES
APPLICANT: GUDAS, JEAN MARIE
TITLE OF INVENTION: MODIFIED PEPTIDES AS THERAPEUTIC AGENTS
FILE REFERENCE: A-527A
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87.0%; Score 20; DB 18; Length 6;
Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2; Indels
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87.0%; Score 20; DB 18; Length 6;
Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
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Gaps
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87.0%; Score 20; DB 19;
Best Local Similarity 60.0%; Pred. No. 4.2e+06;
Matches 3; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                CTHER INFORMATION: Mdm/hdm antagonist peptide US-09-563-286B-130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Search completed: February 12, 2003, 11:24:04 Job time : 100.375 secs
                 PRIOR APPLICATION NUMBER: 09/428,082
PRIOR FILING DATE: 1999-10-22
PRIOR APPLICATION NUMBER: 60/105,371
PRIOR FILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 1151
SOFTWARE: PatentIn version 3.1
CURRENT FILING DATE: 2000-05-03
                                                                                                                                                                                                                TYPE: PRT ORGANISM: Artificial Sequence
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                                                                                                                                                              SEQ ID NO 130
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(without alignments)
59.144 Million cell updates/sec
                                                                                                                                 February 12, 2003, 11:14:12 ; Search time 7.08333 Seconds
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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23
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Pending Patents AA New:*
1: /cgn2 6/ptodata/1/paa/USOG_NEW COMB.pep:*
2: /cgn2 6/ptodata/1/paa/USOG_NEW COMB.pep:*
3: /cgn2 6/ptodata/1/paa/USOB NEW COMB.pep:*
5: /cgn2 6/ptodata/1/paa/USOB NEW COMB.pep:*
5: /cgn2 6/ptodata/1/paa/USOB NEW COMB.pep:*
5: /cgn2 6/ptodata/1/paa/USOB NEW COMB.pep:*
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7: /cgn2 6/ptodata/1/paa/USOB_NEW COMB.pep:*

Maximum Match 100% Listing first 45 summaries

Post-processing: Minimum Match 0%

Database :

seq length: 0 seq length: 25

Minimum DB s Maximum DB s

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

141, App 9, Appli 343, App 199, App 54, Appl 12994, A 15, Appl 12722, A 3, Appli 139, App 149, App 155, App 37, Appl Sequence 286, App Sequence 22, Appl Sequence 27, Appl Sequence 12, Appl Sequence 36, Appl Sequence 36, 7 Sequence 286, Sequence 343, Sequence 149 Description Sequence US-09-701-080C-12 US-10-174-717A-36 US-09-701-080C-22 US-09-701-080C-27 US-06-438-805-9 US-10-211-088-141 US-60-438-805-9 US-10-231-343 US-10-225-567A-2041 US-10-225-567A-2041 US-10-225-567A-209 US-10-225-567A-199 US-10-225-139 US-10-233-599-139 US-10-283-599-139 US-10-283-599-139 US-10-283-599-139 US-10-283-599-139 US-10-283-144-37 US-10-283-144-37 US-10-283-144-37 US-10-171-734-4 US-10-099-056-2094 SUMMARIES Query Match Length Score Result

COMPUTER: IBM Compatible

2094, Ap

Sequence

App

Sequence 1318, Ap	Sequence 316, App	Sequence 399, App	Sequence 13, Appl	Sequence 20, Appl	Sequence 39, Appl	Sequence 31, Appl	Sequence 129, App	Sequence 372, App	Sequence 380, App	Sequence 372, App	Sequence 380, App	Sequence 16, Appl	Sequence 73, Appl	Sequence 73, Appl	Sequence 14, Appl	Seguence 95, Appl	Sequence 251, App	Sequence 7, Appli
US-10-225-567A-1318	PCT-US02-34021-316	PCT-US02-34021-399	US-10-282-958-13	US-10-282-958-20	US-10-282-958-39	PCT-US02-24317-31	US-09-673-166A-129	US-10-062-109A-372	US-10-062-109A-380	US-10-005-480A-372	US-10-005-480A-380	US-09-573-830-16	US-10-062-109A-73	US-10-005-480A-73	US-09-573-830-14	US-10-266-829-95	US-10-133-210-251	US-10-269-171A-7
19 6	20 1	20 1	21 6	21 6	21 6	8	8	9	9	9	9	10 5	10 6	10 6	11 5	14 6	15 6	15 6
8 78.3	8 78.3		8 78.3	8 78.3	8 78.3	7 73.9	7 73.9	7 73.9	7 73.9	7 73.9	7 73.9	7 73.9	7 73.9	7 73.9	7 73.9	7 73.9	7 73.9	7 73.9
27 1	28 1	29 1	30 1	31 1	32 1	33 1	34 1	35 1	36 1	37 1	38 1	39 1	40 1	41 1	42 1	43 1	44	45 1

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Sequence 12, Application US/09701080C
GENERAL INFORMATION:
APPLICANT: INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300 1
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT PILING DATE: 2001-02-27
PRIOR FILING DATE: 1998-05-26
PRIOR FILING DATE: 1998-01-05
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver: 2.1
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Schwegman, Lundberg, Woessner & Kluth P.A. STREET: 121 South Eighth Street, St. 1600
CITY: Minneapolis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87.0%; Score 20; DB 5; Length 7; 60.0%; Pred. No. 3.5e+05;
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                                                                                                                                                                                                                                                                                                                                                                                                   ORGANISM: Artificial Sequence
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ZIP: 55402
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
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US-10-174-717A-36
US-09-701-080C-12
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US-09-701-080C-27
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LENGTH: 13
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LENGTH: 13
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87.0%; Score 20; DB 5; Length 11;
Best Local Similarity 60.0%; Pred. No. 4e+02;
Matches 3; Conservative 0; Mismatches 2; Indels
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APPLICANT Sette, Alessandro
APPLICANT: Sidney, John
APPLICANT: Sidney, John
APPLICANT: Kast, W. Martin
APPLICANT: South Wartin
APPLICANT: Spimmune, Inc.
ITLE OF INVENTION: HLA Binding Peptides and Their Uses
FILE REFERENCE: 39963-20019-20
CURRENT APPLICATION NUMBER: US/09/189, 702A
CURRENT FILING DATE: 1998-11-10
PRIOR APPLICATION NUMBER: US 08/205,713
PRIOR FILING DATE: 1994-03-04
NUMBER OF SEQ ID NOS: 380
SOFTWARE: PASTSEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   87.0%; Score 20; DB 6; Length 7; 60.0%; Pred. No. 3.5e+05; tive 0; Mismatches 2; Indels
             SOFTWARE: FastSEQ Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/10/174,717A
FILING DATE: 18-Jun-2002
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/096,749
FILING DATE: June 12, 1998
APPLICATION NUMBER: 60/049,410
FILING DATE: June 12, 1997
ATTORNEY/ABONT INFORMATION:
NAME: Ann S. VİKSHİNS
REGISTRATION NUMBER: 37,748
REFERENCE/DOCKET NUMBER: 109.034US4
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE: NO
FRAGMENT TYPE: internal
ORIGINAL SOURCE:
SEQUENCE DESCRIPTION: SEQ ID NO: 36:
US-10-174-717A-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; OTHER INFORMATION: p53.14 peptide 34.0324 US-09-189-702A-286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 286, Application US/09189702A GENERAL INFORMATION:
OPERATING SYSTEM: WINDOWS
                                                                                                                                                                                                                                                                                                                                            TELEFAX: (612) 339-3061
                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 7 amino acids
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
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Matches 3; Conserva
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LENGTH: 11
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KENOLICANTI INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPEPTIDES FROM CREB BINDING PROTEIN AND RELATED PROTISIN P300 INTER OF INVENTION: PRANCRIPTIONAL REGULATION
FILE REFERENCE: N73477C GCW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT FILING DATE: 1998-05-26
FRIOR APPLICATION NUMBER: GB 9911303.8
FRIOR FILING DATE: 1998-05-26
FRIOR FILING DATE: 1999-0157.0
FRIOR FILING DATE: 1999-0157.0
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GENERAL INFORMATION:
THOUSE THAT INSTITUTE OF MOLECULAR AND CELL BIOLOGY
TITLE OF INVENTION: POLYPERTIDES FROM CREB BINDING PROTEIN AND RELATED PROTEIN P300
TITLE OF INVENTION: TRANSCRIPTIONAL REGULATION
FILE REFERENCE: N7347C GCW
CURRENT APPLICATION NUMBER: US/09/701,080C
CURRENT FILING DATE: 2001-02-27
PRIOR APPLICATION NUMBER: GB 9811303.8
PRIOR PLICATION NUMBER: GB 990157.0
PRIOR PLILNG DATE: 1998-05-26
PRIOR PLILNG DATE: 1999-01-05
NUMBER OF SEQ ID NOS: 36
SOFTHARE: PATCHILN UNEL 1005.3
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APPLICANT: Bright, Gary R.
APPLICANT: Premkumar, D. David
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Best Local Similarity 60.v.
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US-10-203-138A-12722
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87.0%; Score 20; DB 6; I
Best Local Similarity 60.0%; Pred. No. 6.9e+02;
Matches 3; Conservative 0; Mismatches 2;
                                                                                                          87.0%; Score 20; DB 1; I
60.0%; Pred. No. 6.4e+02;
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                                                                                                                                            0; Mismatches
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GENERAL INFORMATION:
APPLICANT: Baxter, John
APPLICANT: Darimont, Beatrice
; APPLICANT: Feng, Weijun
                                  NAME/KEY: MISC FEATURE
)
OTHER INFORMATION: DGI7-20M-4-G12
PCT-US02-34021-343
                                                                                                                                              3; Conservative
   ORGANISM: Eschericia coli
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                                                                                                                           Best Local Similarity
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US-10-203-138A-12722
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                                                                                                                Query Match
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Segence 343, Application PC/TUS0234021
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: DGI BioTechnologies, Inc.
TITLE OF INVENTION: TARGET SPECIFIC SCREENING AND ITS USE FOR IDENTIFYING TARGET BIND
FILE REFERENCE: 2598-4009PC
CURRENT APPLICATION NUMBER: PCT/US02/34021
CURRENT PILING DATE: 2002-10-24
PRIOR FILING DATE: 2001-10-24
PRIOR FILING DATE: 2001-10-24
NUMBER OF SEQ ID NOS: 537
SOFTWARE: PARCHIN Version 3.1
SEQ ID NO 343
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APPLICANT: Chen, Yih-Tai
TITLE OF INVENTION: Novel Fusion Proteins And Assays For Molecular Binding
FILE REFERENCE: 01-1022-10
FILE REFERENCE: 01-1022-10
CURRENT APPLICATION NUMBER: US/10/211,088
CURRENT FILING DATE: 2002-10-15
PRIOR APPLICATION NUMBER: 60/309,395
PRIOR PILING DATE: 2001-08-01
PRIOR PLING DATE: 2001-08-01
PRIOR PLING DATE: 2001-12-13
NUMBER OF SEQ ID NOS: 366
SOFTWARE: Patentin version 3.1
SEQ ID NO 141
LENGTH: 15
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US-60-438-805-9
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GENERAL INFORMATION:
APPLICANT: KODADEK, THOMAS;
TITLE OF INVENTION: MIXED ELEMENT CAPTURE ACENTS;
TITLE OF INVENTION: MIXED ELEMENT CAPTURE ACENTS;
CURRENT APPLICATION NUMBER: US/60/438,805;
CURRENT FILING DATE: 2003-01-09;
NUMBER OF SEQ ID NOS: 12;
SOFTWARE: Patentin Ver. 2.1;
SEQ ID NO 9
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ORGANISM: Artificial Sequence
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US-60-438-805-9
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APPLICANT: LifeSpan Biosciences
APPLICANT: Brown, Joseph P.
APPLICANT: Brown, Joseph P.
APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Glenna C.
APPLICANT: Burmer, Glenna C.
APPLICANT: Roush, Christine L.
TITLE OF INVENTION: ANYIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
FILE REPERRNCE: 1920-4-4
CURRENT APPLICATION NUMBER: US/10/225,567A
PRIOR TELLING DATE: 2000-12-19
PRIOR FILING DATE: 2000-12-19
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GENERAL INFORMATION:
APPLICANT: Dasseux, Jean-Louis
APPLICANT: Sekul, Renate
APPLICANT: Cornut, Isabelle
APPLICANT: Gornut, Isabelle
APPLICANT: Gornut, Isabelle
APPLICANT: Metz, Gunthar
TITLE OF INVENTION: GUBPLY APOLIPOPROTEIN A-I AGONISTS AND THEIR
TITLE OF INVENTION: USE TO TREAT DYSLIPIDEMIC DISORDERS.
NUMBER OF SEQUENCES: 274
CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds LLP
STREET: 115 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                       Gaps
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FILE REFERENCE: 1920-4-4
CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT FILING DATE: 2001-12-19
FRIOR PRILING DATE: 2000-12-19
NUMBER OF SEQ ID NOS: 2292
SOFTWARE: Patentin version 3.1
SSOFTWARE: Patentin version 3.1

LENGTH: 14
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SOFTWARE: PatentIn version 3.1
SEQ ID NO 1920
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Best Local Similarity 60.09
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CRGANISM: Homo sapiens
US-10-225-567A-2196
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Best Local Similarity
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US-10-225-567A-1920
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US-10-283-599-199
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APPLICANT: Brown, Joseph P.
APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Glenna C.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORE
FILE REFERENCE: 1200-4-4
CURRENT APPLICATION NUMBER: US/10/225,567A
CURRENT PILING DATE: 2000-12-19
PRIOR APPLICATION NUMBER: 60/257,144
NUMBER OF SEQ ID NOS: 2292
SOFTWARE: PatentIn version 3.1
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GENERAL INFORMATION:
APPLICANT: Brown, Joseph P.
APPLICANT: Burmer, Glenna C.
APPLICANT: Burmer, Glenna C.
APPLICANT: Roush, Christine L.
TITLE OF INVENTION: ANTIGENIC PEPTIDES AND ANTIBODIES FOR G PROTEIN-COUPLED RECEPTORS
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                                                  APPLICANT: Wagner, Richard
APPLICANT: Yamamoto, Keith
TITLE OF INVENTION: METHODS AND COMPOUNDS FOR MODULATING NUCLEAR RECEPTOR
TITLE OF INVENTION: COACTIVAOR BINDING
FILE REFREENCE: 9811-008-599
CURRENT FILING DATE: 1999-03.30
CURRENT FILING DATE: 1999-03.30
PRIOR APPLICATION NUMBER: US 60/079,956
PRIOR PILING DATE: 1998-03.30
NUMBER OF SEQ ID NOS: 60
SOFTWARE: Patentin version 3.0
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; OTHER INFORMATION: Xaa is any amino acid US-09-281-717A-3
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Fletterick, Robert
                     Kushner, Peter
West, Brian
Wagner, Richard
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 3; Conserv
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APPLICANT:
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APPLICANT:
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Search completed: February 12, 2003, 11:24:44 Job time: 7.08333 secs
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TOPOLOGY: linear
MOLECULE TYPE: peptide
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US-10-319-786-54
; Sequence 54, Application US/10319786
; Sequence 54, Application US/10319786
; GENERAL INFORMATION:
; APPLICANT: Little, Roger G
; TITLE OF INVENTION: Biologically Active Peptides from
; TITLE OF PROFINITY PROFICE TO THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF THE PROFICE OF T
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STREET: 10 South Wacker Drive, Suite 3000
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REGISTRATION NUMBER: 30,742
REFERENCE/DOCKET NUMBER: 009196-0007-999
RELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELERAX: 650-493-4935
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 199:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 amino acids
ZIF: 10036-2811
COMPUTER READBLIB FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FASLSEQ VERSION 2.0
CURRENT APPLICATION DATA:
FILING DATE: 29-OCT-2002
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/940,136
FILING DATE: 29-SEP-1997
ATTORNEY/AGENT INFORMATION:
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COMPUTER READABLE FORM:
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STATE: Illinois
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
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              FILING DATE: 11-MAR-1995
ATTORNEY AGENT INCRMATION:
NAME: McDonnell, John J.
REGISTRATION NUMBER: 26,949
REFRENCE/DOCKET NUMBER: 93,1133-J
TELEPHONE: 312-715-1000
TELEFAX: 312-715-1000
TELEFAX: 312-715-1000
TELEFAX: 312-715-1000
SEQUENCE CHARACTER: 512-51
APPLICATION NUMBER: US 08/209,762
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                                                                    February 12, 2003, 11:12:42; Search time 4.58333 Seconds (without alignments) 32.098 Million cell updates/sec
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Sequence 15,
Sequence 19,
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Sequence 29,
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Sequence 3,
Sequence 6,
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Sequence 3,
Sequence 6,
Sequence 1,
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Sequence 1
Sequence 3
Sequence 2
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/cgm2 6/ptodata/1/iaa/5B COMB.pep:*
/cgm2 6/ptodata/1/iaa/6A COMB.pep:*
/cgm2 6/ptodata/1/iaa/6B COMB.pep:*
/cgm2 6/ptodata/1/iaa/B CTWS.COMB.pep:*
/cgm2 6/ptodata/1/iaa/PcTWS.COMB.pep:*
GenCore version 5.1.3
Copyright (c) 1993 - 2003 Compugen Ltd.
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US-08-424-957-29

US-08-277-660A-2

US-08-277-660A-3

US-08-424-957-2

US-08-424-957-2

US-08-424-957-3

US-08-424-957-3

US-08-424-957-3

US-09-035-686-2

US-09-035-686-3

US-09-035-686-3

US-09-035-686-3

US-09-035-686-3

US-09-035-686-1

US-08-424-957-15

US-08-426-17-18

US-08-426-17-14

US-08-709-173-11

US-08-709-173-11
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Maximum Match 100%
Listing first 45 summaries
                                                OM protein - protein search, using sw model
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Match Length DB
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Maximum DB seq length: 25
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91.3%; Score 21; DB 1; Length 11; 60.0%; Pred. No. 1.5e+02; ative 0; Mismatches 2; Indels

Conservative

Query Match Best Local Similarity

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STRANDEDNESS: TOPOLOGY: linear amino acid

; US-08-277-660A-16

ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter 1, 190
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-60244/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids

21,	equence 8,	Ξ;		,	80	11,	12,	18,	11	12	Sequence 18, Appl	ν,	07.	177	12,	equence 13,								٠		acion inereci		Horbort	2														
US-08-638-911A-21	-08-709-177	-08-709-177-1	-709-177-1	US-08-277-660A-7	US-08-277-660A-8	US-08-424-957-11	US-08-424-957-12	US-08-424-957-18	US-09-035-686-11	US-09-035-686-12	US-09-035-686-18	-08-277-660A-9	-08-277-660A-1	-08-277-660A	-08-277-660A	-08-277-66	US-08-277-660A-15	ALIGNMENTS		277660A			т.	•	Interruption of Binding of MDMZ	n and Therapeutic Application		4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4 4	Suite 3400							ble	-Dos	se #1.0, Version #1.30		US/08/277,660A			
6	9 2				,	-	-	Н	4	4	4		11 1							Sequence 16, Application US/08277660A			ley, Steven M	å			ES: 27	Dresse:	ADDRESSES: Flent, nombacin, lest,		1000	United States	7	FORM:	MEDIUM TYPE: Floppy disk	~		Patentin Release	ON DATA:			514	. NOTHANGORNI TNGOR/ VGNGOTHK
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20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20	20		RESULT 1	nce 16,	It No. 57	GENERAL INFORMATION	APPLICANT: Picksley,	APPLICANT:	TITLE OF INVENTION:	TITLE OF INVENTION:	NOMBER OF SEQUENCES: Z	RESPOND	AUDKESSE CTUTETT.	OTTAC:		>	ZIP: 94111-4187	COMPUTER READABLE FORM:	MEDIUM 1	COMPUTER	OPERATING SYSTEM:	SOFTWARE:	CURRENT APPLICATION DATA:	APPLICATION NUMBER:	FILING DATE:	CLASSIFICATION:	2/VONDOTIO
28	29	30	31	32	33	34	35	36	37	38	39	40	41	42	43	44	45		RESULT	Seque;	, Paten	GENE	; AF	; AF	; TI	TI	 N		••							٠.			บ 	.,			

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                                                                       Sequence 29, Application US/08424957
Patent No. 5770377
Batent No. 5770377

APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
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CITY: San Francisco
STREET: Four Embarcadero Center, Suite 3400
COUNTRY: United States
COUNTRY: IMP COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING STREET: INP PC COMPATING
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Patent No. 6153391

GENERAL INFORMATION:

APPLICANT: David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof

CORRESPONDENCE ADDRESS:

CORRESPONDENCE ADDRESS:

ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert

STREET: Four Embarcadero Center, Suite 3400

STATE: California

COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 91.3%; Score 21; DB 1; Length 11; Best Local Similarity 60.0%; Pred. No. 1.5e+02; Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
CLASSIFICATION: DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGRY INCRMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: 1.1.00
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
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94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
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RESULT 2
US-08-424-957-29
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Gaps
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APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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91.3%; Score 21; DB 4; Length 11;
Best Local Similarity 60.0%; Pred. No. 1.5e+02;
Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Flehr, Hohbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERALING STAIRS:

OCHARALING STAIRS:

CURRENT APPLICATION DATA:

APPLICATION NUMBER:

TILING DATE:

CLASSIFICATION:

ATTORNEY/AGENT INFORMATION:

NAME:

REGISTRATION NUMBER:

AC044/WHD

TELECOMMUNICATION NUMBER:

TELECOMMUNICATION NUMBER:

TELECOMMUNICATION NUMBER:

TELECOMMUNICATION INPORMATION:

TELEFRAX: (415) 398-3249

TELEFRAX: 910 277299
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPBRATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: DEGGEY, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: (415) 398-3249
TELEFEX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
LENGTH: 11 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; Sequence 2, Application US/08277660A; Patent No. 5702908
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy dif
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          unknown
                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 FXXLW 5
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US-09-035-686-29
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1 FXXLW 5
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                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Dicksley, Steven M.
APPLICANT: Lane, David M.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
ADDRESSEB: Flehr, Hobbach, Test, Albritton & Herbert STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .;
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                                                                                                                                                               2; Indels
                                                                                                                          87.0%; Score 20; DB 1; Length 6; 60.0%; Pred. No. 2e+05; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A-60244/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE: C. C. C. C. CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H. REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-602
TELECOMMUNICATION INFORMATION: (415) 398-3249
TELERA: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                   Sequence 3, Application US/08277660A
Patent No. 5702908
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 6
US-08-424-957-2
; Sequence 2, Application US/08424957
; Patent No. 57-0377
; GENERAL INPORMATION:
; APPLICANT: PICKSley, Steven M.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: United States
ZIP: 94111-4187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 6 amino acids
SEQUENCE CHARACTERISTICS
                         LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid STRANDEDNESS:
                                        TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FOPOLOGY: linear
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Matches 3; Conserv
                                                                              linear
                                                                                                                              Query Match
Best Local Similarity
Matches 3; Conserv
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                                                                                                                                                                                                  1 FXXLW 5
                                                                                                                                                                                                                                    2 FSDLW 6
                                                                            ; TOPOLOGY:
US-08-277-660A-2
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US-08-277-660A-3
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APPLICANT: Picksley, Steven M.
APPLICANT: Dane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof NUMBER OF EXQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          87.0%; Score 20, DB 1; Length 6; 60.0%; Pred. No. 2e+05; 2; Indels ative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: California COMPUTEY: United States
ZIP: 94111-4187
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPAtible
COMPOTER: IBM PC COMPAtible
COMPUTER: IBM PC COMPATIBLE
COMPOTER: IBM PC COMPATIBLE
COMPOTER: IBM PC COMPATIBLE
SOFTWARE: PATENTIN Release #1.0, Version #1.30
SOFTWARE: PATENTIN DATA:
APPLICATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTONEY/AGENT INFORMATION:
MANNEY ATTONEY/AGENT INFORMATION:
                   PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOWMUNICATION INFORMATION:
TELEFAX: (415) 398-3249
TELEFXX: (415) 398-3249
TELEFXX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 3:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-424-957-6; Sequence 6, Application US/08424957; Patent No. 5770377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEPHONE: (415) 781-1989
TELERAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 60.0
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: unknown
US-08-424-957-3
                                                                                                                                                                                                                                                                                                                                                                                           TYPE: amino acid STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYFE: amino acid STRANDEDNESS:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 FXXLW 5
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US-09-035-686-2
Sequence 2, Application US/09035686
Patent No. 6153991
Sequence 2, Application US/09035686
Patent No. 6153991
Patent No. 6153991
TREAL INFORMATION:
APPLICANT: Pickeley, Steven M.
APPLICANT: Lame, David P.
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Gaps
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Query Match 87.0%; Score 20; DB 1; Length 6; Best Local Similarity 60.0%; Pred. No. 2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          87.0%; Score 20; DB 4; Length 6; 60.0%; Pred. No. 2e+05; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: Patentin Release #1.0, Version #1.30 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
FILING DATE: 19-APR-1995
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERRACE/DOCKET NUMBER: 34,190
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 10
US-09-035-686-3
; Sequence 3, Application US/09035686
; Patent No. 6153391
; GENERAL INFORMATION:
; APPLICANT: Picksley, Steven M.
; APPLICANT: Lane, David P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 60.0
Matches 3; Conservative
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TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: unknown US-09-035-686-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
                                                                                        1 FXXLW 5
                                                                                                                                  2 FSELW 6
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Patent No. 6153391

GENERAL INFORMATION:

APPLICANT: Lane, David P.

TITLE OF INVENTION: Interruption of Binding of MDM2 and P53

TITLE OF INVENTION: Protein and Therapeutic Application Thereof

NUMBER OF SEQUENCES: 50

CORRESPONDENCE ADDRESS:

ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert

STREET: Four Embarcadero Center, Suite 3400

CITY: San Francisco

STATE: California

COUNTRY: United States

COUNTRY: United States

COMPUTER IBM PC Compatible

COMPUTER: IBM PC Compatible

COMPUTER: IBM PC Compatible

COMPUTER: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Gaps
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
ADDRESSER: Flahr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 87.0%; Score 20; DB 4; Length 6; 60.0%; Pred. No. 2e+05; tive 0; Mismatches 2; Indels
                                                                                                                                                    CITY: San Francisco
CITY: San Francisco
CITY: California
COUNTRY: United States
ZIP: 94111-4187
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: BY PC compatible
COMPUTER: BY PC compatible
COMPUTER: BY PC compatible
COMPUTER: STATEM: PC-DOS/MS-DOS
SOFTWARE: PAPELICATION DATA:
APPLICATION NUMBER: US/09/035,686
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REGISTRANCE/DOCKET NUMBER: A-61228/WHD
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIPICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-APR-1995
APPLICATION UNDERE: US 08/277,660
FILING DATE: 20-JUL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 3: SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 60.0
Matches 3; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: amino acid
STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                               FILING DATE
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APPLICANT: Kaelin, William
APPLICANT: Jost, Christine
TITLE OF INVENTION: METHODS OF TREATMENT USING
TITLE OF INVENTION: NBS-1, ANTIBODIES AND PROTEINS THERETO, AND USES OF THE
TITLE OF INVENTION: ANTIBODIES
TITLE OF INVENTION: ANTIBODIES
CORRESPONDENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 87.0%; Score 20; DB 4; Length 6; Best Local Similarity 60.0%; Pred. No. 2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows
SOFTWARE: FASELSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/081,975
FILING DATE: 12-MAY-1998
CLASSIFICATION A435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/046,207
FILING DATE: 12-MAY-1997
ATTORNEY/AGENT INFORMATION:
             CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/424,957
FILING DATE: 19-ARR-1995
ATTORNEY/ACBNT INFORMATION:
NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61228/WHD
TELEPHONE: (415) 781-1999
TELEPHONE: (415) 781-1999
TELEPAX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                    A-61228/WHD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 12
US-09-081-975-1
; Sequence 1, Application US/09081975
; Patent No. 6451979
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Nixon Peabody LLP
STREET: 101 Federal Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      30,628
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Eisenstein, Ronald I
REGISTRATION NUMBER: 30,628
REFERENCE/DOCKET NUMBER: 47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 6 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 617-345-1300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compati
                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: unknown
                                                                                                                                                                                                                                                                                                                                                                                                    amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: US
ZIP: 02110
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TOPOLOGY: unknown US-08-424-957-15
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STRANDEDNESS:
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                                                                 ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
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                                                                                                                                            0; Gaps
                                                                                                                                                                                                                                                                                                                   Sequence 27, Application US/08277660A
Patent No. 5702908
GENERAL INFORMATION
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ;
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Patent No. 5770377
GENERAL INFORMATION:
APPLICANT: Picksley, Steven M.
APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
                                                                                               Query Match 87.0%; Score 20; DB 4; Length 6; Best Local Similarity 60.0%; Pred. No. 2e+05; Matches 3; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              87.0%; Score 20; DB 1; Length 7; 60.0%; Pred. No. 2e+05; Live 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert
STREET: Four Embarcadero Center, Suite 3400
CITY: San Francisco
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: United States
ZIP: 94111-4187
ZORPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-60244/WHD
TELECOMOUNICATION INFORMATION:
TELEPHONE: (415) 781-1989
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/277,660A
FILING DATE: 20-JUL-1994
CLASSIFICATION: 514
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (415) 398-3249
TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 27:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           : 7 amino acids amino acid
TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 California
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Best Local Similarity
'2.2 3; Conserve
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US-08-277-660A-27
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                                                                                                                                                                                                                        2 FSDLW 6
                                                                                                                                                                                                                                                                                                    US-08-277-660A-27
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US-08-424-957-15
                                                US-09-081-975-1
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Gaps
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APPLICANT: Lane, David P.
TITLE OF INVENTION: Interruption of Binding of MDM2 and P53
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
TITLE OF INVENTION: Protein and Therapeutic Application Thereof
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                                                                   E: Flehr, Hohbach, Test, Albritton & Herbert
Four Embarcadero Center, Suite 3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Flehr, Hobbach, Test, Albritton & Herbert STRET: Pour Embarcadero Center, Suite 3400 CITY: San Francisco STATE: California
                                                                                                                                                                                                                                                                                                                            SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
PLACATION NUMBER: US/08/424,957
FILING DATE: 19-APR-1995
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APPLICATION NUMBER: US/08/424,957
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A-61228/WHD
                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE: 19-APR-1995
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/277,660
FILING DATE: 20-ULL-1994
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMpatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 19, Application US/08424957; Patent No. 5770377
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Dreger, Walter H.
REGISTRATION NUMBER: 24,190
REFERENCE/DOCKET NUMBER: A-61;
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 398-3249
TELEFAX: (415) 398-3249
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: 910 277299
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                       COUNTRY: United States ZIP: 94111-4187
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                  NUMBER OF SEQUENCES: 5: CORRESPONDENCE ADDRESS:
                                                                                   STREET: Four Languages CITY: San Francisco GTATE: California GTATE: California Stat
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy of
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PRIOR APPLICATION DATA:
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### APPLICATION NUMBER: US 08/277,660
### FILING DATE: 20-JUL-1994
### ATTONEY AGENT INPORMATION:
### NAME: Dreger, Walter H.
### REGISTRATION NUMBER: 24,190
### REFERENCE/DOCKET NUMBER: 24,190
### REFERENCE/DOCKET NUMBER: A-61228/WHD
### TELEPHONE: (415) 398-3249
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